Supporting Information

Lipid Membrane Interaction of Peptide/DNA Complexes Designed for Gene Delivery

*Neval Yilmaz*¹, *Yutaka Kodama*^{1,2} *and Keiji Numata*^{1,3*}

¹Biomacromolecules Research Team, RIKEN Center for Sustainable Resource Science, Wako, Saitama, Japan

²Center for Bioscience Research & Education, Utsunomiya University, Tochigi, Japan

³Laboratory for Biomaterial Chemistry, Department of Material Chemistry, Graduate School of Engineering, Kyoto University, Japan

*e-mail: numata.keiji.3n@kyoto-u.ac.jp

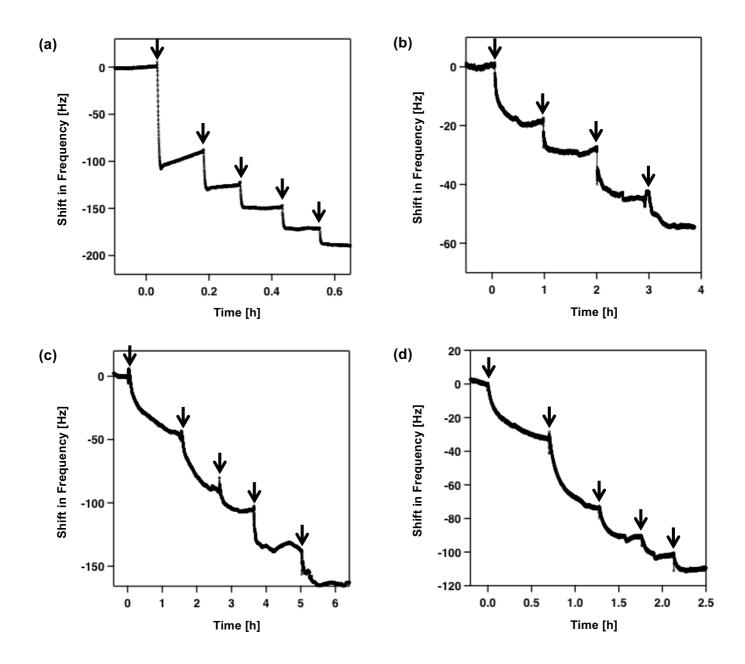


Figure S1. Time-course plots showing the binding of Cytcox (a), BP100 (b), Cytcox-KH₉ (c), and BP100-KH₉ (d) to Plasma-MM. The arrows indicate the injection of the peptide. The concentrations of the injected amounts of Cytcox, BP100, Cytcox-KH₉, and BP100-KH₉ were 41, 2.2, 1.3, and 0.7 μ M, respectively.

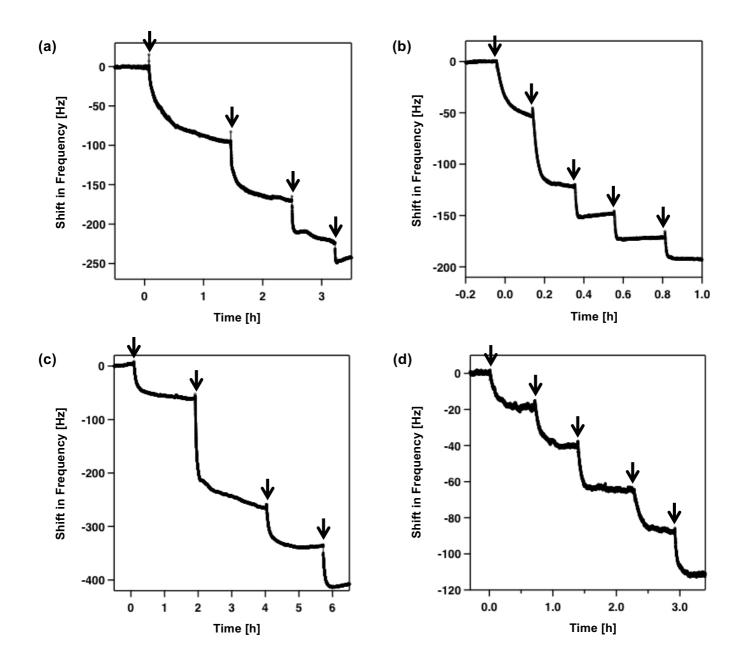


Figure S2. Time-course plots showing the binding of Cytcox (a), BP100 (b), Cytcox-KH₉ (c), and BP100-KH₉ (d) to Mito-MM. The arrows indicate the injection of the peptide. The concentrations of the injected amounts of Cytcox, BP100, Cytcox-KH₉, and BP100-KH₉ were 8.3, 9.1, 1.3, and 0.7 μ M, respectively.

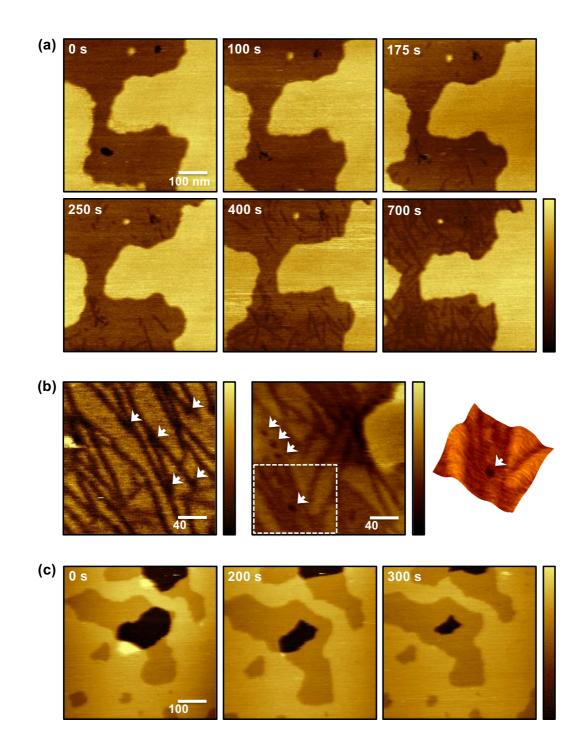


Figure S3. HS-AFM height images showing the changes in Plasma-MM in the presence of 8 μ M Cytcox (**a**, **b**) and 9 μ M BP100 (**c**). "0 s" in A and C correspond to approximately 5 and 9 min after addition of Cytcox and BP100, respectively. The arrows in B point to the pore-like defects. The 3D image is the region in the dashed line box. The color bar indicates the Z-range between 0 (darkest) to Z_{max} (brightest); $Z_{max}(a) = 9$ nm, $Z_{max}(b) = 2.5$ nm (left), 6 nm (right), and $Z_{max}(c) = 10$ nm.

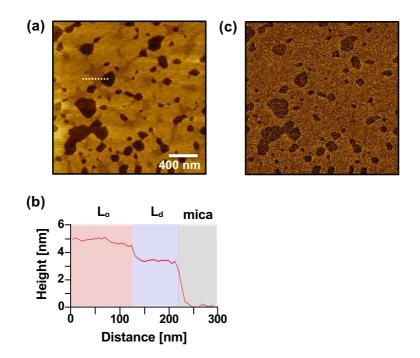


Figure S4. HS-AFM height (a) and phase (c) images showing the height and phase variation in plasma-MM. (b) Height profile along the dashed line in (a).

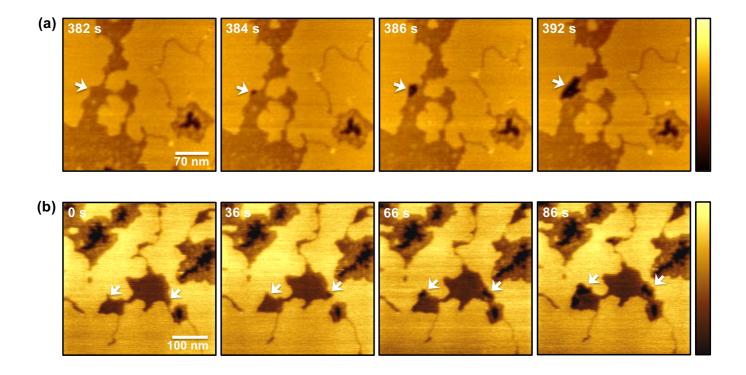


Figure S5. HS-AFM height images showing the pore formation in Plasma-MM in the presence of 3 μ M Cytcox-KH₉ (**a**) and 0.5 μ M BP100-KH₉ (**b**). "0 s" in (b) corresponds to 28 min after addition of BP100-KH₉ and 26 min after the start of observation in Figure 4b. The arrows point to the pores. The color bar indicates the Z-range between 0 (darkest) to Z_{max} (brightest); Z_{max}(**a**, **b**) = 5 nm.

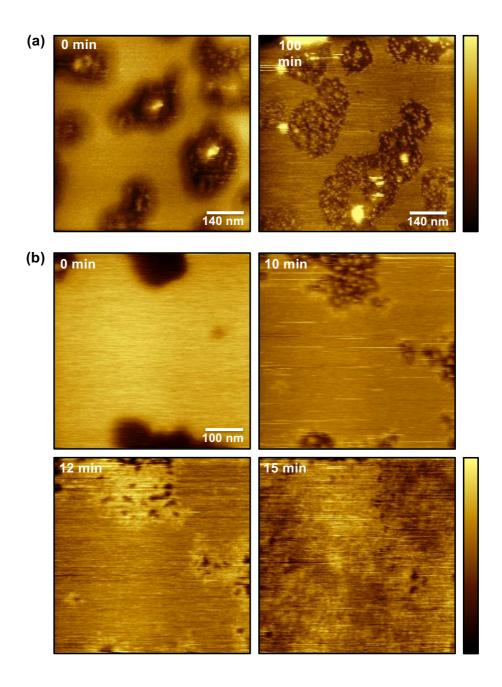


Figure S6. HS-AFM height images showing the changes in Mito-MM in the presence of 12 μ M Cytcox (a) and 1.3 μ M Cytcox-KH₉ (b). "0 min" corresponds to the time just after addition of peptide. The color bar indicates the Z-range between 0 (darkest) to Z_{max} (brightest); $Z_{max}(a) = 7$ nm and $Z_{max}(b) = 6$ nm (0 min, 10 min), 4 nm (12 min), 3 nm (15 min).

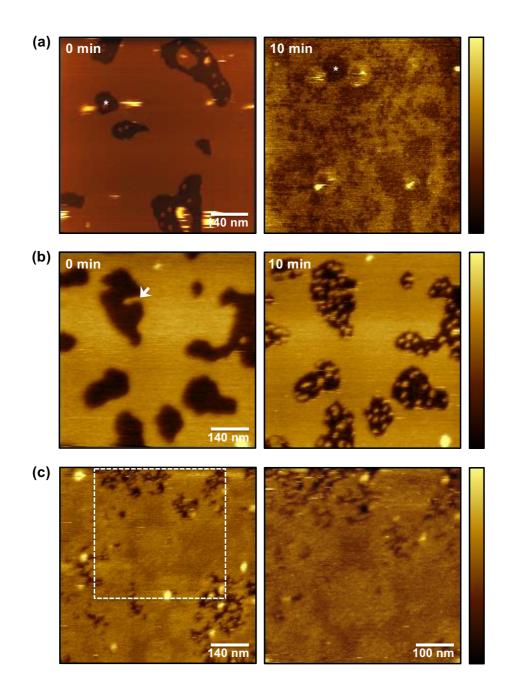


Figure S7. HS-AFM height images showing the changes in Mito-MM in the presence of 9 μ M BP100 (a) and 0.4 μ M BP100-KH₉ (b, c). "0 min" corresponds to the time just after addition of peptide. The arrow in (b) points to the tubule-like structure protruding from the membrane in the presence of BP100-KH₉. The images in (c) show the thin layer formed on Mito-MM 45 min after addition of BP100-KH₉. The image to the right is the higher-resolution image of the region in the dashed line box. The color bar indicates the Z-range between 0 (darkest) to Z_{max} (brightest); $Z_{max}(a) = 40$ nm (0 min), 5 nm (10 min), $Z_{max}(b) = 12$ nm, and $Z_{max}(c) = 10$ nm.

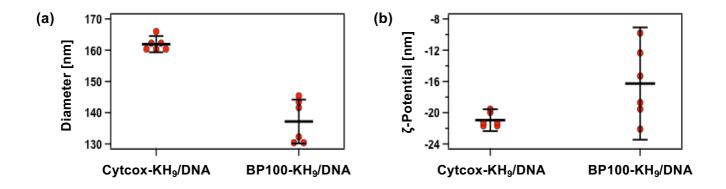


Figure S8. Hydrodynamic size (a) and ζ-potential (b) of Cytcox-KH₉/DNA and BP100-KH₉/DNA complexes measured by DLS. The number of data points was 6 for each measurement. The highest and lowest lines indicate the standard deviation from the mean value (centered line).

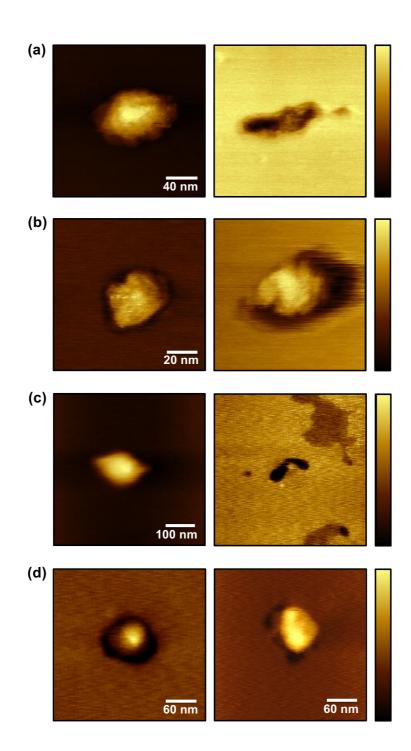
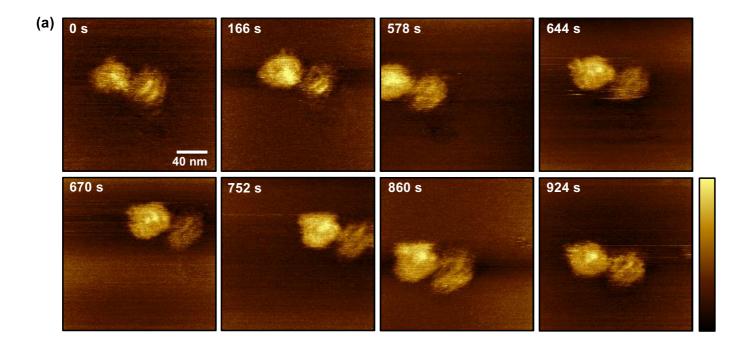


Figure S9. HS-AFM height images showing the complexes of Cytcox-KH₉/DNA (**a**-**c**) and BP100-KH₉/DNA (**d**) on Plasma-MM. The images to the right in (a)-(c) were recorded at a higher imaging force to reveal the disrupted region by Cytcox-KH₉/DNA complex. The images in (d) show the disrupted regions of the membrane around BP100-KH₉/DNA complex. The color bar indicates the Z-range between 0 (darkest) to Z_{max} (brightest); $Z_{max}(a) = 22$ nm (left), 6 nm (right), $Z_{max}(b) = 6$ nm (left), 8 nm (right), $Z_{max}(c) = 40$ nm (left), 6 nm (right), and $Z_{max}(d) = 13$ nm.



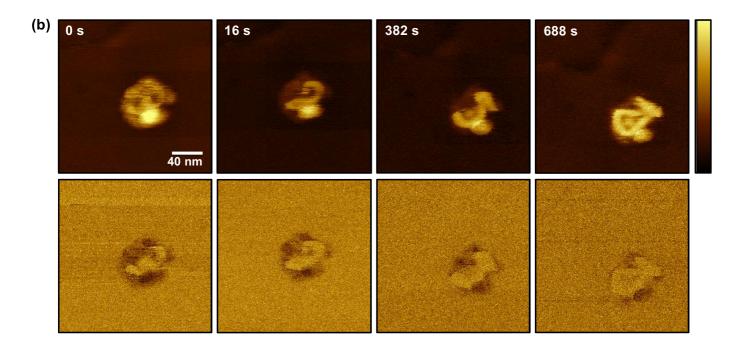


Figure S10. HS-AFM height images of Cytcox-KH₉/DNA complex on Plasma-MM (**a**, **b**) at pH 5.8 in 20 mM MES containing 150 mM NaCl. The phase images in the lower row in (b) show the DNA-rich region of the complex on the peptide-rich darker region. The color bar indicates the Z-range between 0 (darkest) to Z_{max} (brightest); $Z_{max}(a) = 10$ nm and $Z_{max}(b) = 8$ nm.

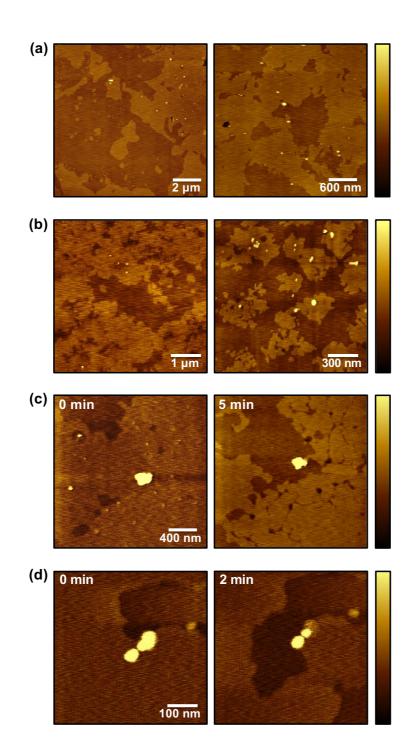


Figure S11. HS-AFM height images showing the localization of Cytcox-KH₉/DNA complex (**a**) and BP100-KH₉/DNA complex (**b**) on L_o domains of Plasma-MM and fluidization of L_o domains by Cytcox-KH₉/DNA complex (**c**) and BP100-KH₉/DNA complex (**d**). The color bar indicates the Z-range between 0 (darkest) to Z_{max} (brightest); $Z_{max}(a) = 15$ nm, $Z_{max}(b) = 10$ nm, $Z_{max}(c) = 7.5$ nm (0 min), 10 nm (5 min), and $Z_{max}(d) = 7$ nm.

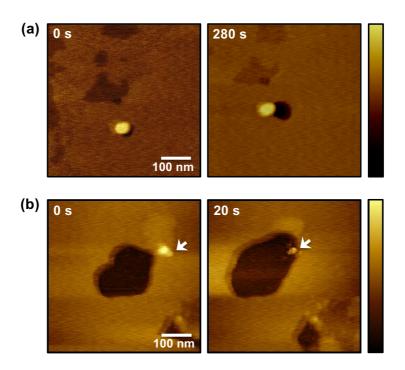


Figure S12. HS-AFM height images showing the pore formation by Cytcox-KH₉/DNA complex in Plasma-MM (**a**, **b**). The color bar indicates the Z-range between 0 (darkest) to Z_{max} (brightest); $Z_{max}(a) = 10 \text{ nm} (0 \text{ s}), 14 \text{ nm} (280 \text{ s}) \text{ and } Z_{max}(b) = 14 \text{ nm}.$

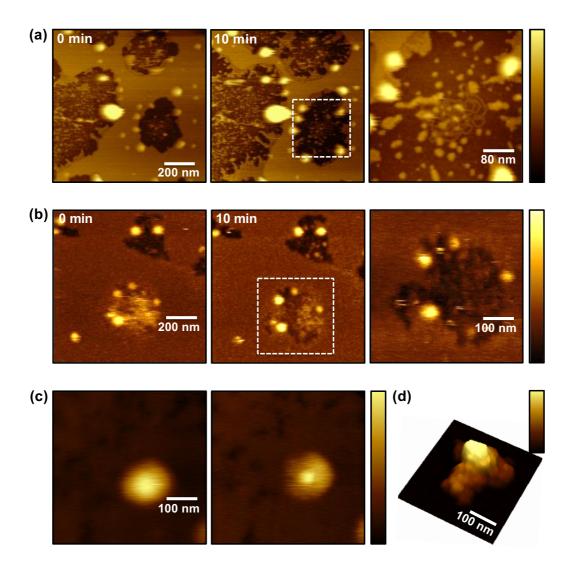


Figure S13. HS-AFM height images showing the disruption in Mito-MM caused by Cytcox-KH₉/DNA complex (a) and BP100-KH₉/DNA complex (b). "0 min" corresponds to 4 h in a and b after addition of 40 μ L of complex into the imaging medium. The third images in A and B are the higher-resolution images of the regions within the dashed line boxes in the second images. The images in (c) and (d) show the intact complex and the partially-dissociated complex on the membrane. The image to the right in (c) was recorded at a higher imaging force. The color bar indicates the Z-range between 0 (darkest) to Z_{max} (brightest); $Z_{max}(a) = 20$ nm, $Z_{max}(b) = 25$ nm, $Z_{max}(c) = 32$ nm (left), 25 nm (right), and $Z_{max}(d) = 100$ nm.

Movies

Movie 1. Cytcox-induced changes in Plasma-MM. Scan rate was 0.5 frame/s.

Movie 2. BP100-induced changes in Plasma-MM. Scan rate was 0.5 frame/s.

Movie 3. Cytcox-KH₉-induced changes in Plasma-MM. Scan rate was 0.5 frame/s.

Movie 4. BP100-KH₉-induced changes in Plasma-MM. Scan rate was 0.5 frame/s.

Movie 5. BP100-KH₉-induced changes in Plasma-MM. Scan rate was 0.5 frame/s. (continuation of Movie 2)

Movie 6. Cytcox-KH₉-induced changes in Mito-MM. Scan rate was 0.5 frame/s.

Movie 7. BP100-KH₉-induced changes in Mito-MM. Scan rate was 0.5 frame/s.