

# Literature Report

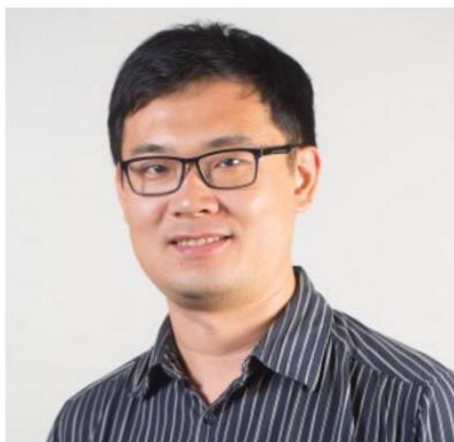
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**Reporter: Liu Weiwei**  
**Date: 2021-01-07**

## Efficient and selective DNA modification on bacterial membranes

Qian Tian,<sup>a</sup> Yousef Bagheri,<sup>a</sup> Puspam Keshri,<sup>a</sup> Rigumula Wu,<sup>a</sup> Kewei Ren,<sup>a</sup> Qikun Yu,<sup>a</sup> Bin Zhao,<sup>a</sup> and Mingxu You<sup>\*a</sup>

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**Dr. Mingxu You**

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- B.S. degree in Chemistry from Peking University in 2008
- Ph.D. degree in Chemistry from the University of Florida in 2012, developing DNA-based devices for cancer diagnosis, targeted drug delivery, and cell membrane biophysical studies.
- Prior to join UMass, he pursued his postdoctoral research with Prof. Samie R. Jaffrey at Weill Medical College of Cornell University, interested in developing RNA-based fluorescent sensors for imaging metabolites and signaling molecules in live cells

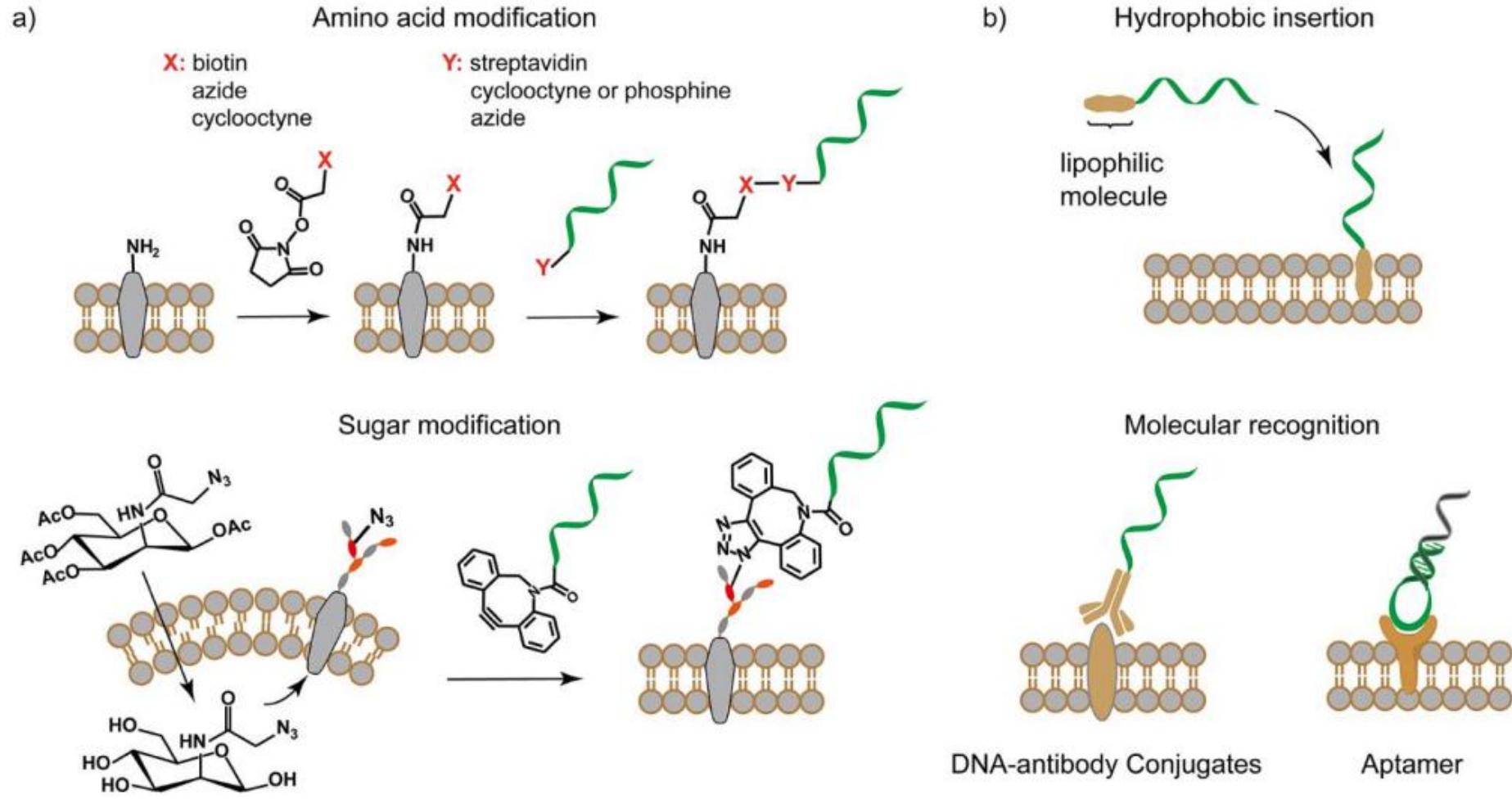
- (1) Biosensors and Bioimaging
- (2) Cell Membrane Biophysics
- (3) DNA/RNA Nanotechnology
- (4) SELEX
- (5) Photo-controlled Devices



# Introduction



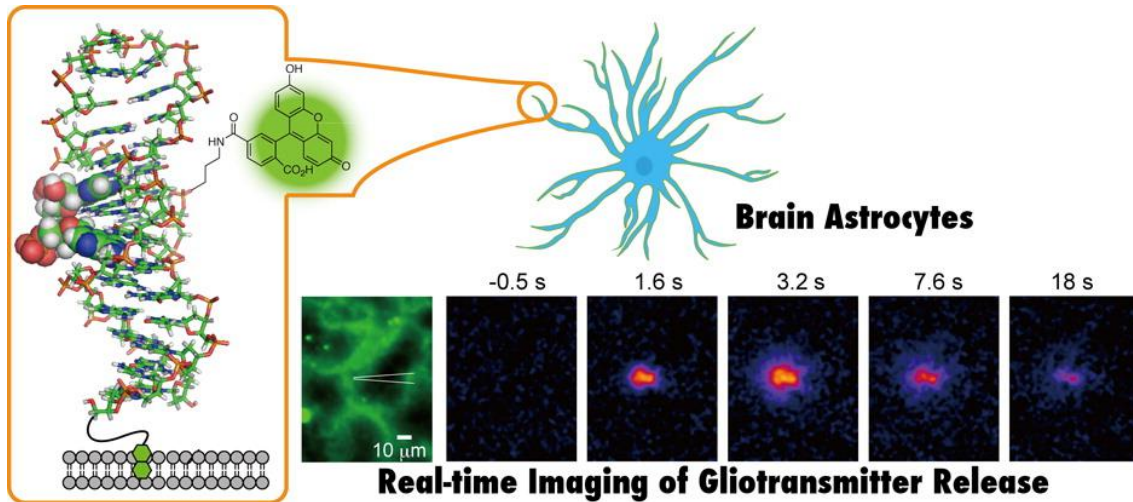
## Synthetic DNA for Cell Surface Engineering



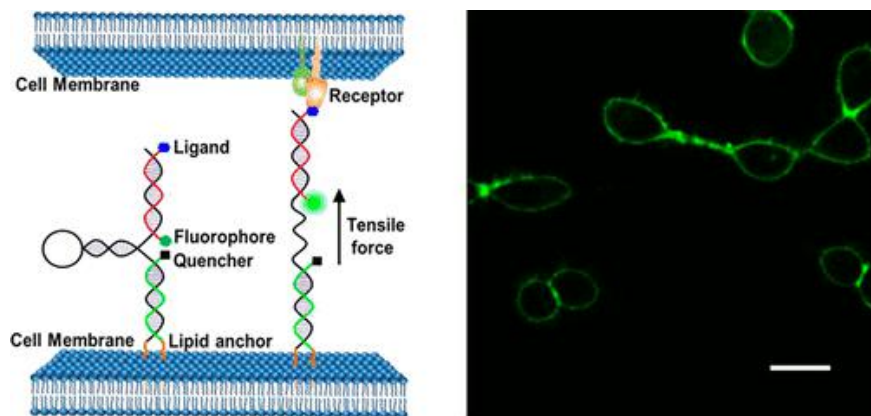
# Introduction



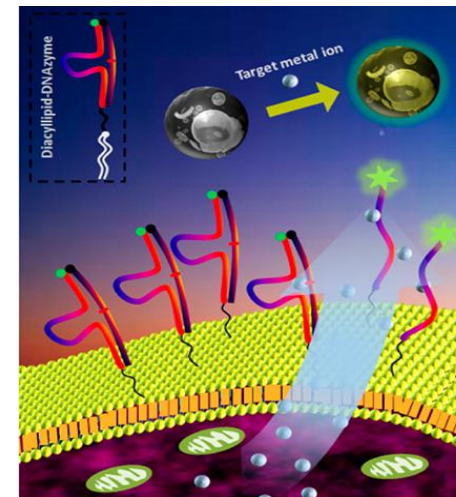
Lipid-DNA conjugates for cell membrane analysis



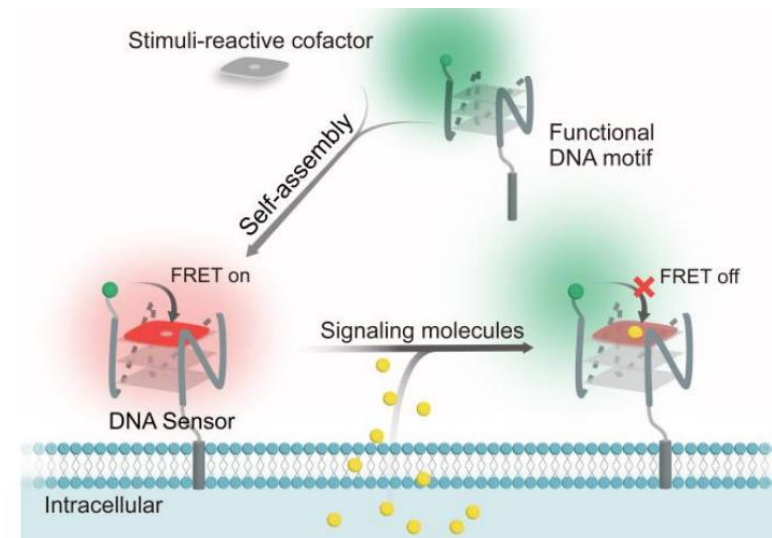
J. Am. Chem. Soc. 2012, 134, 23, 9561–9564



J. Am. Chem. Soc. 2017, 139, 50, 18182



J. Am. Chem. Soc. 2014, 136, 13090-13093

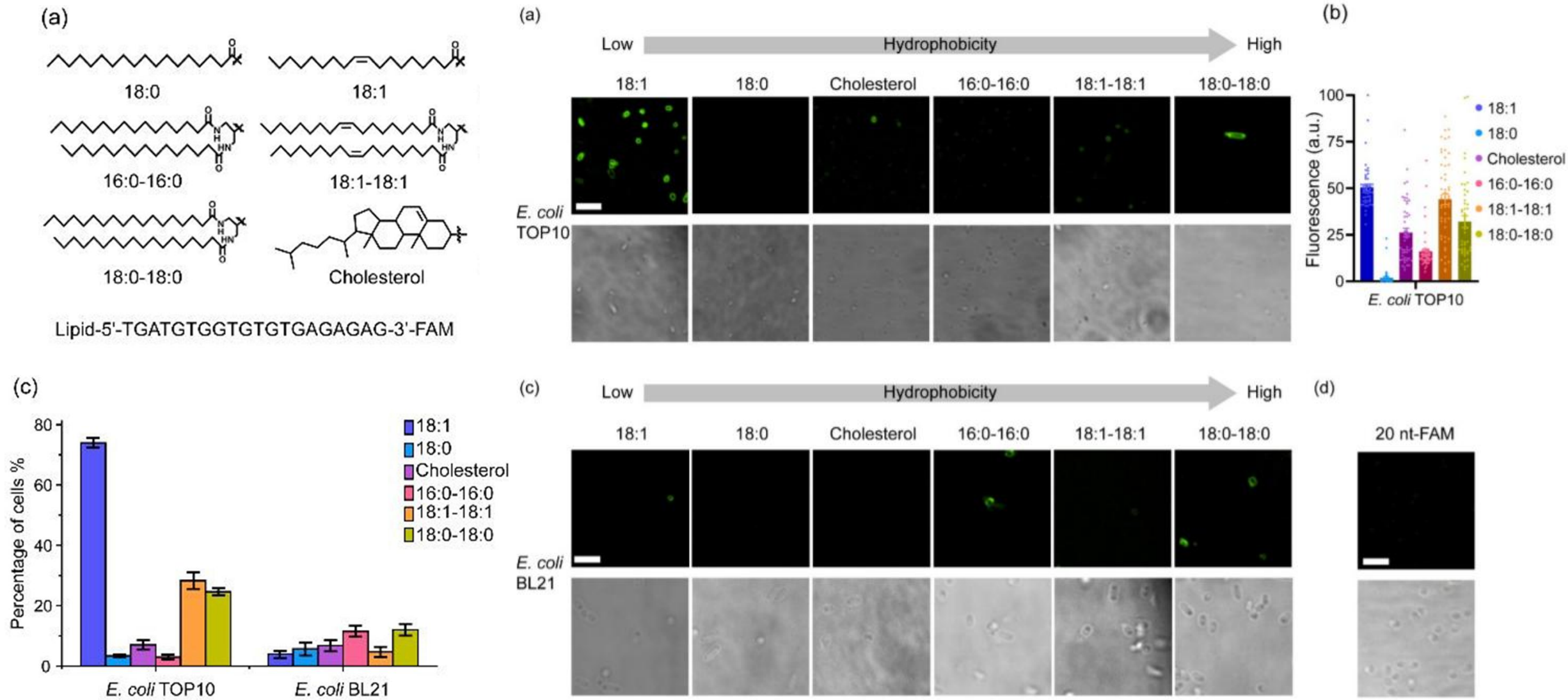


Angew. Chem. Int. Ed. 10.1002/anie.201901320





# Design and bacterial membrane insertion of lipid-DNA conjugates



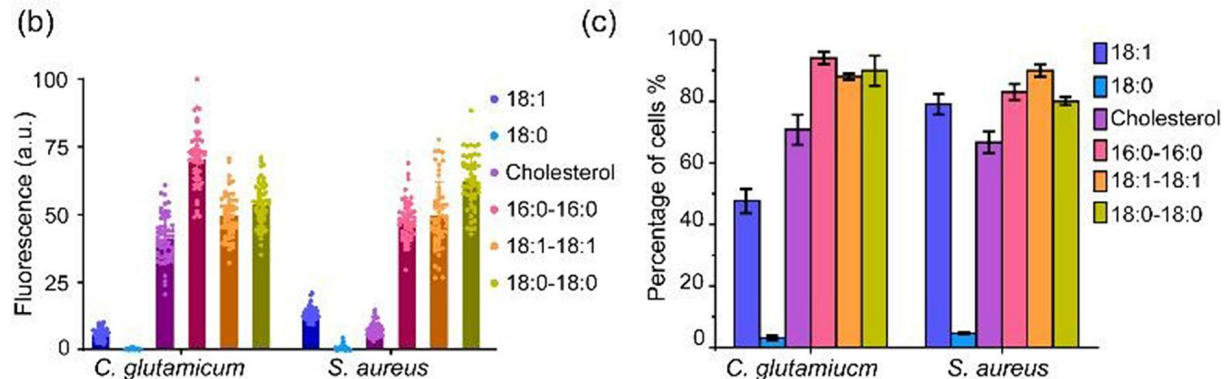
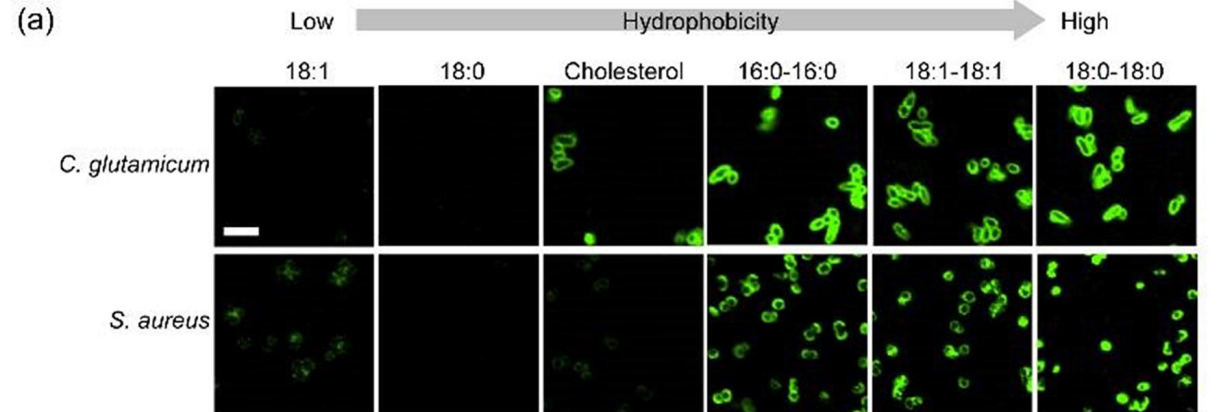
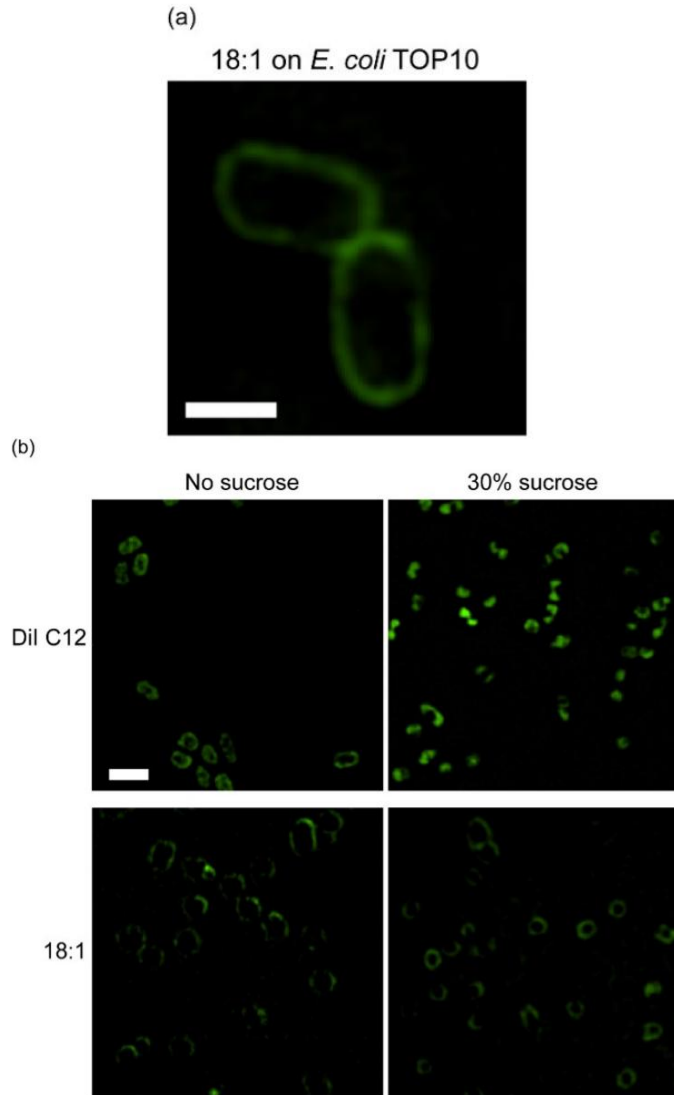


# Design and bacterial membrane insertion of lipid-DNA conjugates

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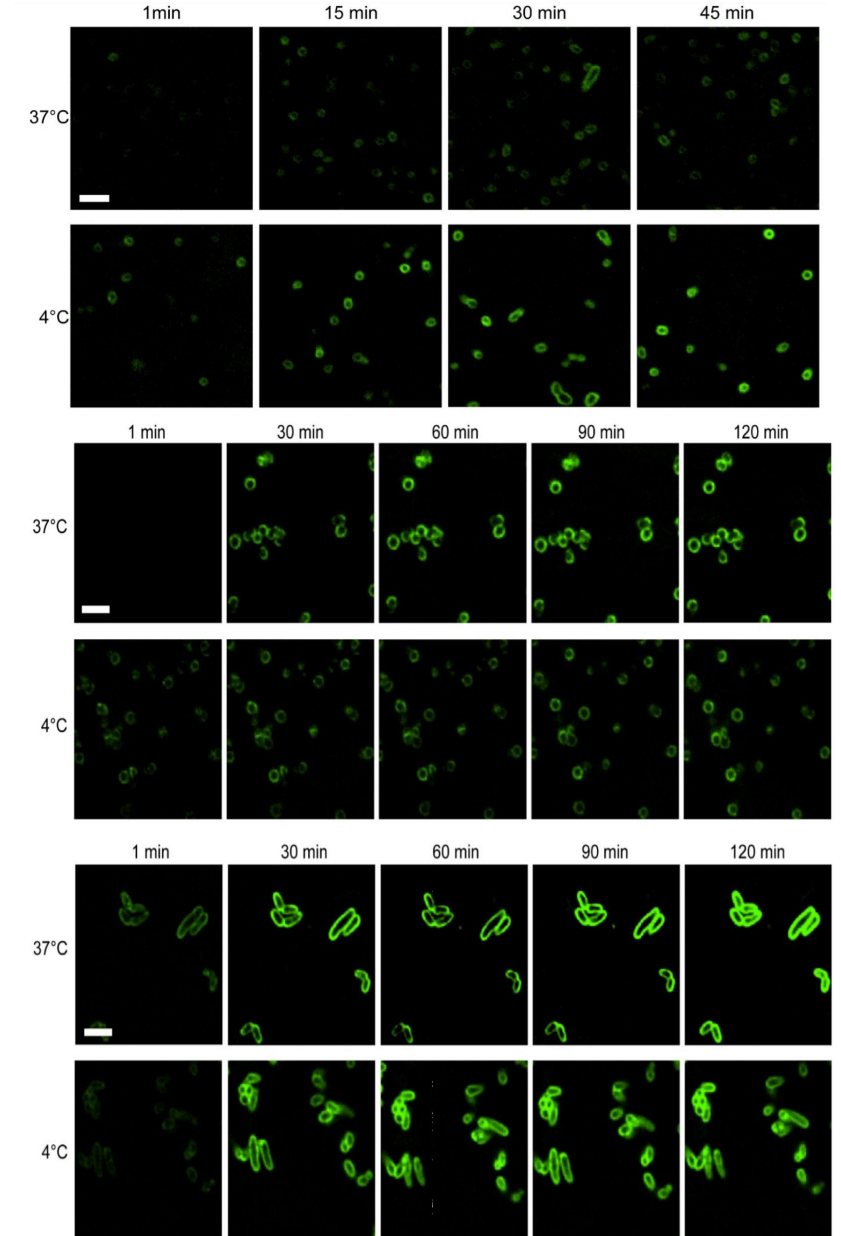
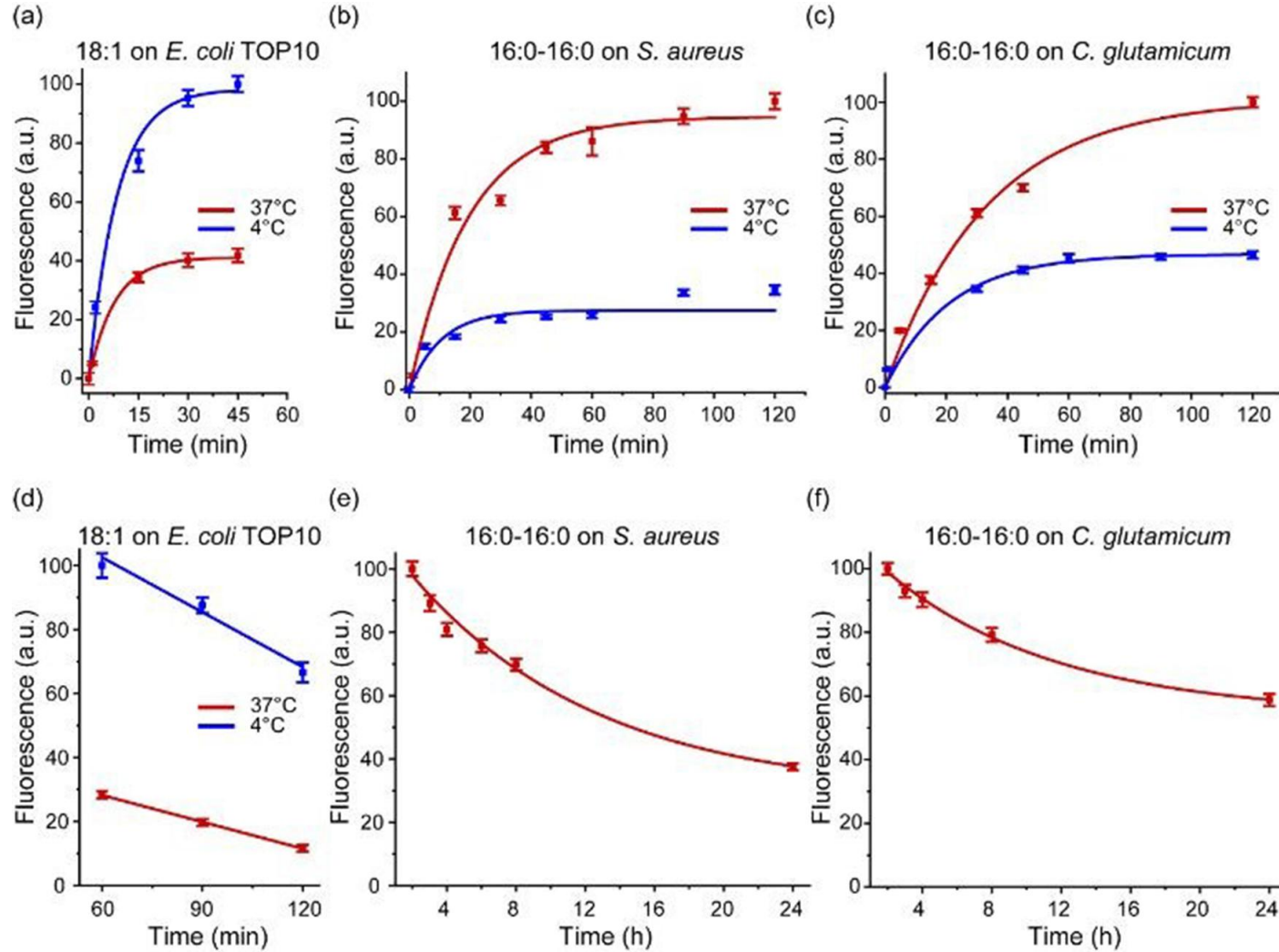
Table S2. Bacterial membrane modification percentage of each lipid-DNA conjugate.

	<i>E. coli</i> TOP10	<i>E. coli</i> BL21	<i>P. aeruginosa</i>	<i>C. glutamicum</i>	<i>S. aureus</i>	<i>M. luteus</i>
18:1	84 ± 4.4	8.1 ± 0.9	<1.0	52 ± 5.5	87 ± 2.8	5.4 ± 0.5
18:0	3.9 ± 0.6	7.0 ± 1.4	<1.0	4.7 ± 0.8	5.5 ± 1.7	<1.0
Cholesterol	9.9 ± 0.9	10 ± 2.3	<1.0	71 ± 1.1	77 ± 3.7	11 ± 1.8
16:0-16:0	4.8 ± 0.4	13 ± 1.3	<1.0	94 ± 2.9	83 ± 2.1	<1.0
18:1-18:1	31 ± 1.9	6.7 ± 1.2	<1.0	88 ± 2.0	90 ± 1.2	8.8 ± 2.8
18:0-18:0	33 ± 3.2	17 ± 1.6	<1.0	90 ± 1.4	80 ± 5.2	<1.0





# Membrane insertion kinetics & persistence of lipid-DNA conjugates

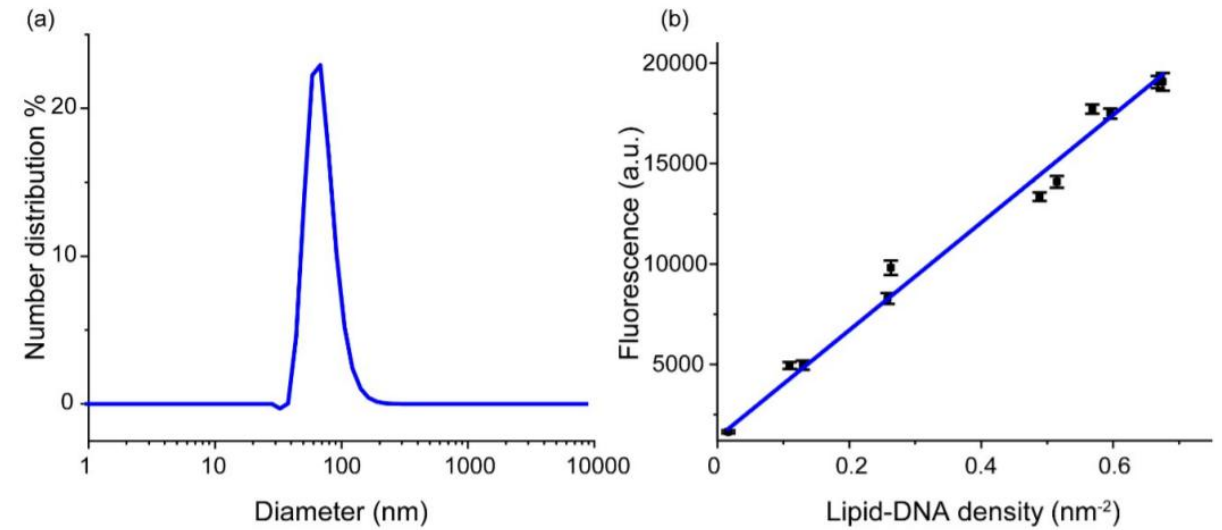
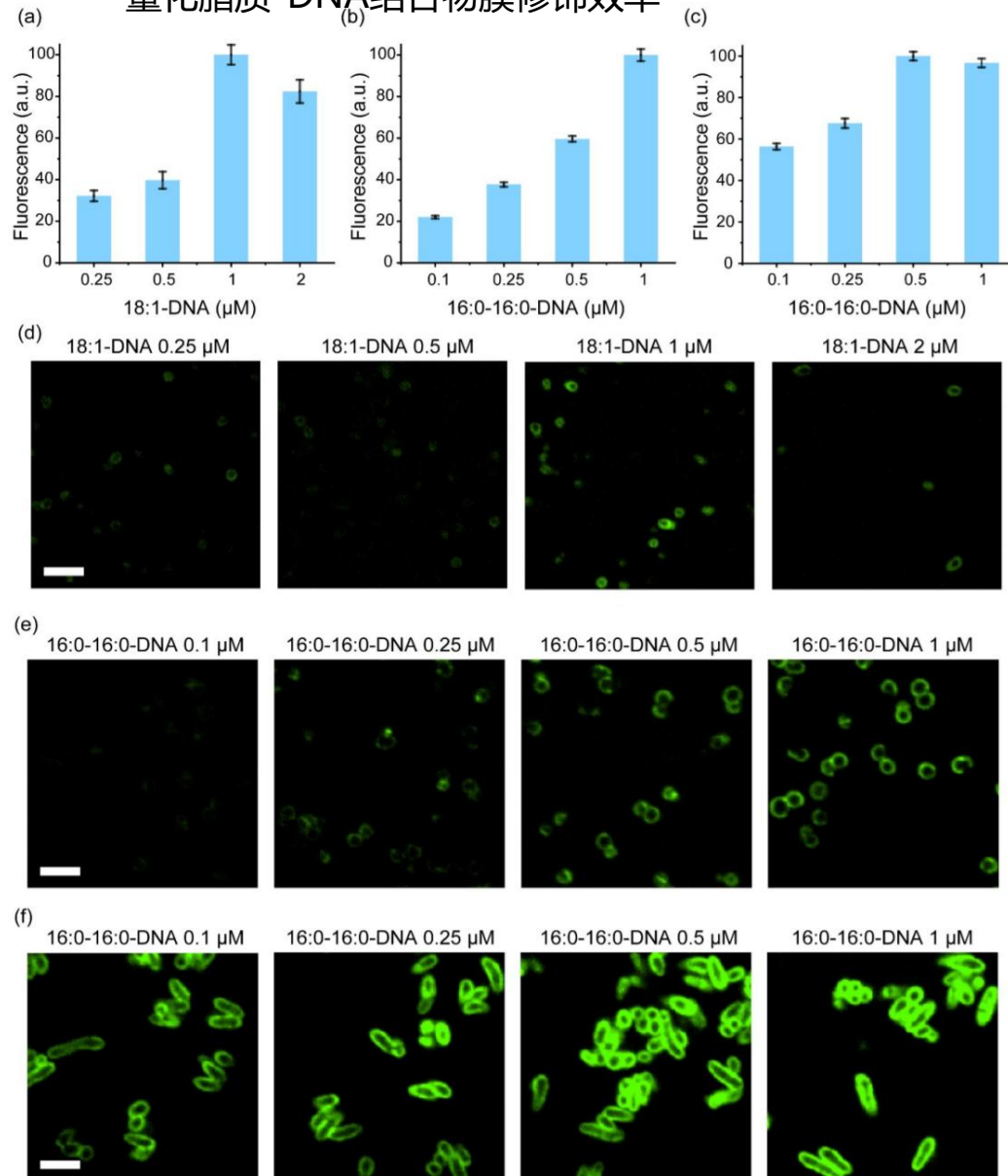






# Membrane modification efficiency of lipid-DNA conjugates

量化脂质-DNA结合物膜修饰效率



**Table S3.** Maximum bacterial membrane density of each lipid-DNA conjugate (unit:  $\text{nm}^{-2}$ ).

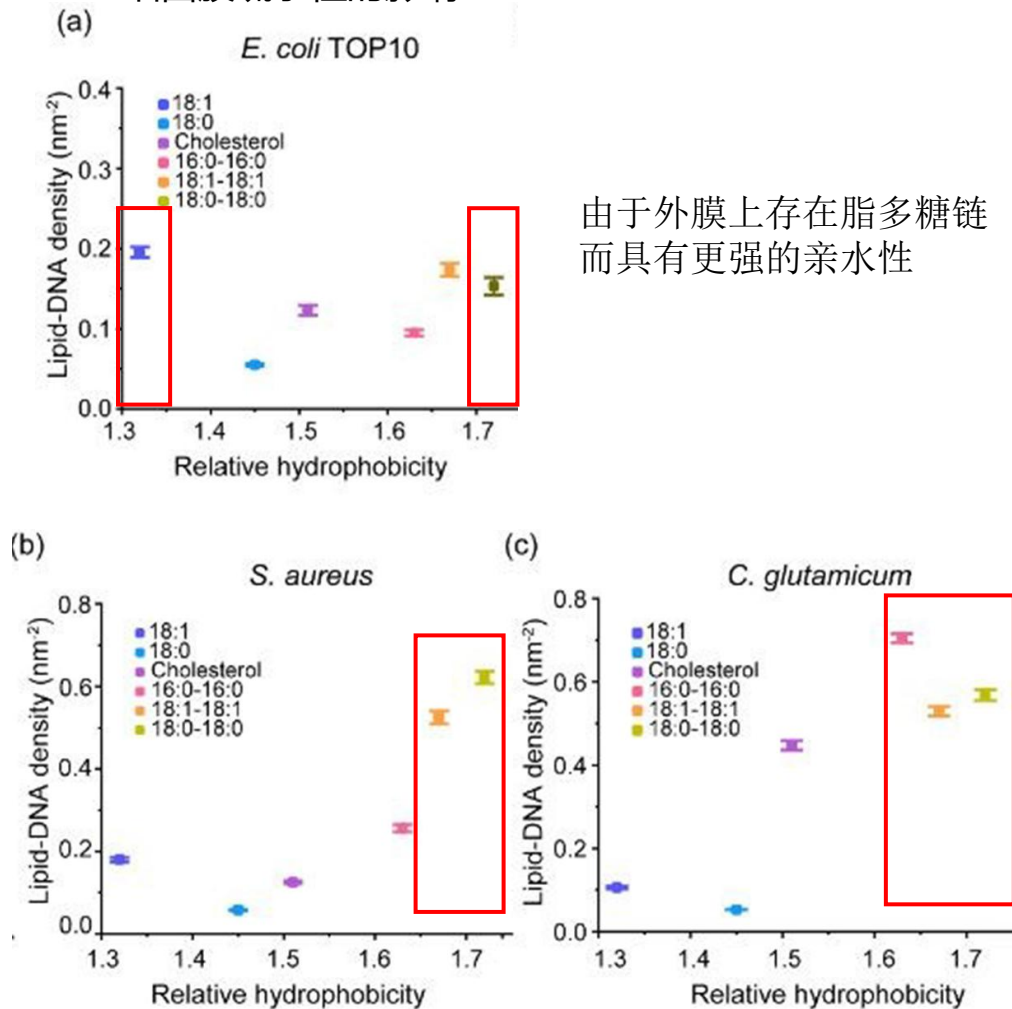
	18:1	18:0	Cholesterol	16:0-16:0	18:1-18:1	18:0-18:0
<i>E. coli</i> TOP10	$0.20 \pm 0.01$	$0.06 \pm 0.01$	$0.12 \pm 0.01$	$0.10 \pm 0.01$	$0.17 \pm 0.01$	$0.15 \pm 0.01$
<i>S. aureus</i>	$0.18 \pm 0.01$	$0.06 \pm 0.01$	$0.13 \pm 0.01$	$0.26 \pm 0.01$	$0.53 \pm 0.02$	$0.62 \pm 0.02$
<i>C. glutamicum</i>	$0.11 \pm 0.01$	$0.05 \pm 0.01$	$0.45 \pm 0.01$	$0.71 \pm 0.01$	$0.53 \pm 0.10$	$0.57 \pm 0.01$



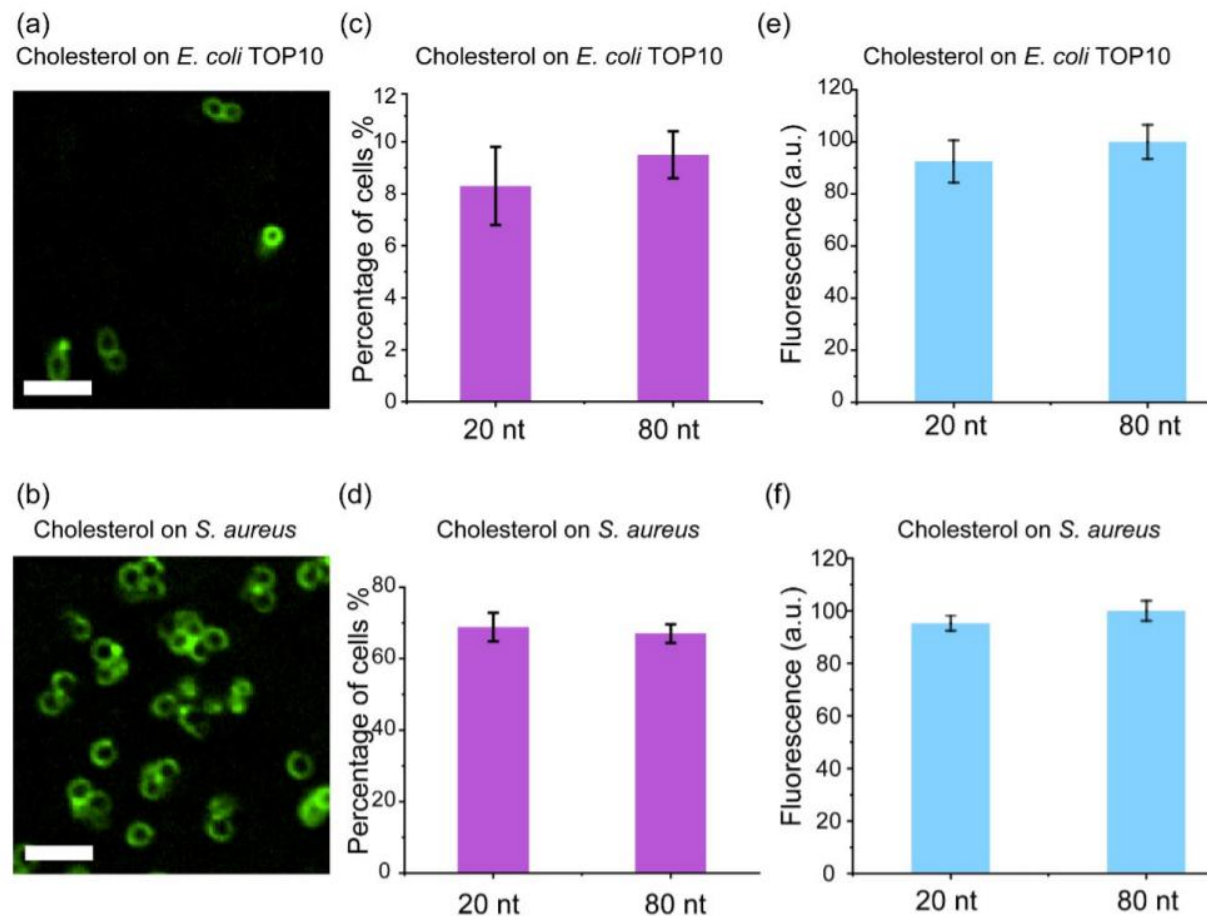


# Effect of lipid-DNA hydrophobicity on the membrane insertion

## 细菌膜疏水性的影响



## DNA长度的影响

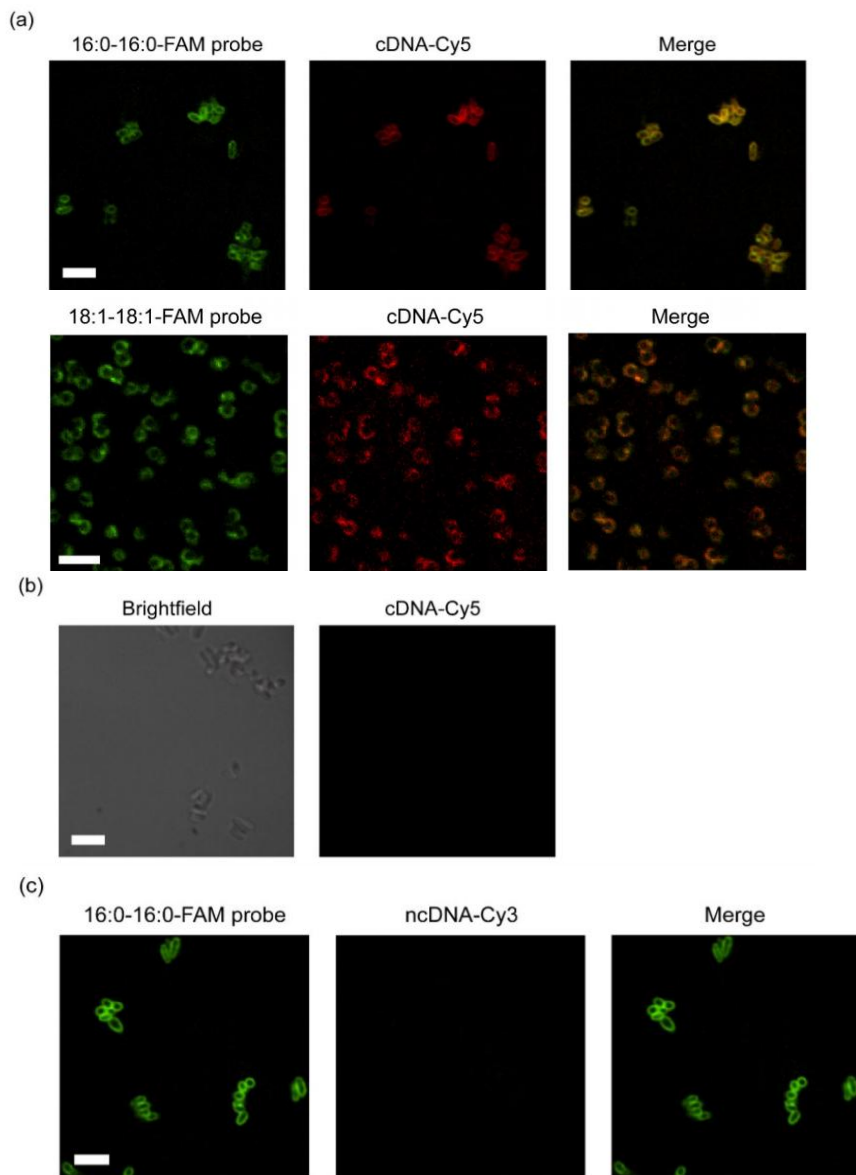


膜含有大量高度疏水的支链氨基酸和脂肪酸

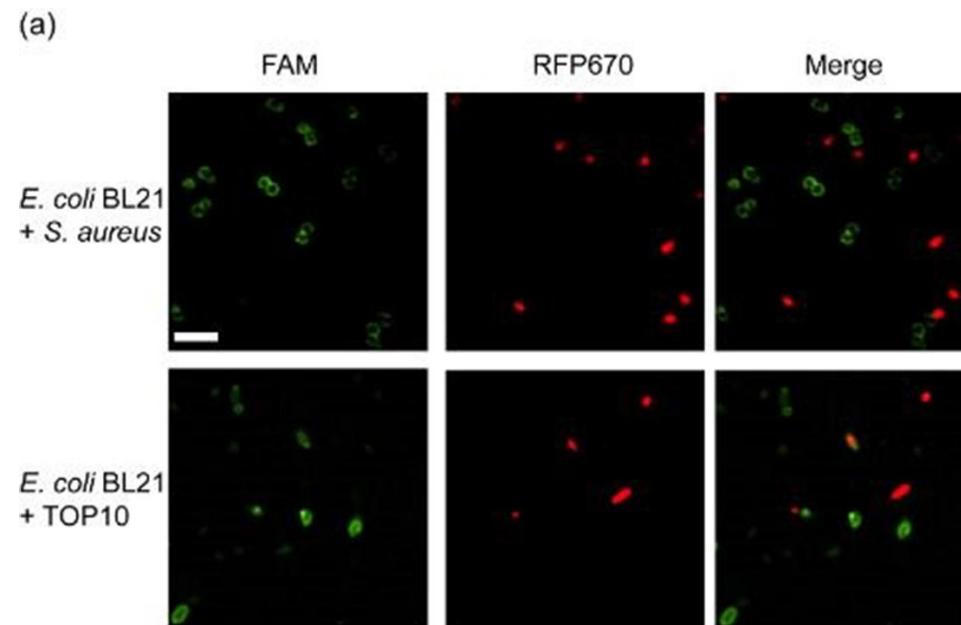


# Selective targeting and detection of bacteria

## DNA杂交



## 检测混合菌液中的细菌

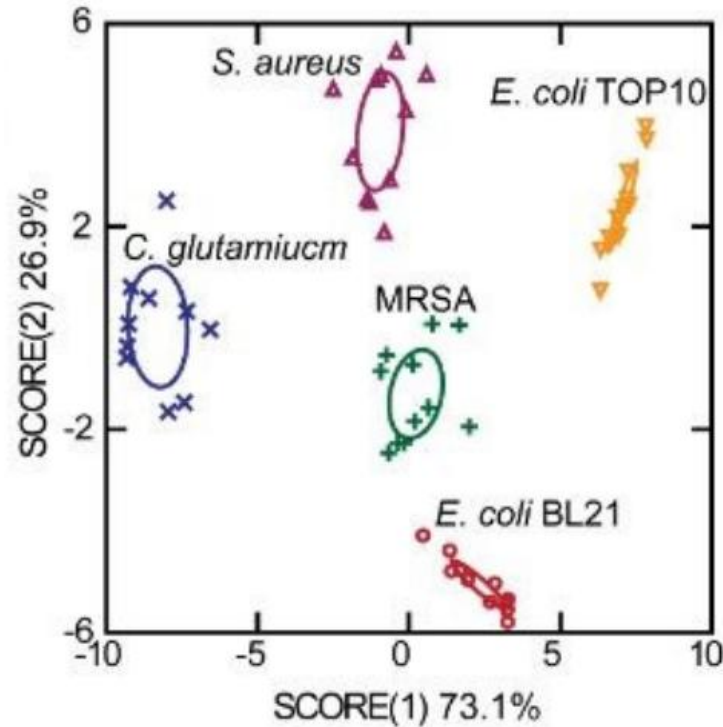


**Figure 5a.** (Top) FAM-labeled 18:1-18:1-DNA conjugate can be used to distinguish *S. aureus* cells from a mixture with RFP670-expressing *E. coli* BL21 cells. (Bottom) Similarly, FAM-labeled 18:1-DNA conjugate was used to distinguish *E. coli* TOP10 cells from a mixture with BL21. Here, 1  $\mu$ M of the lipid-DNA conjugate was incubated with the cell mixture for 1 h at 37°C. Scale bar, 5  $\mu$ m.

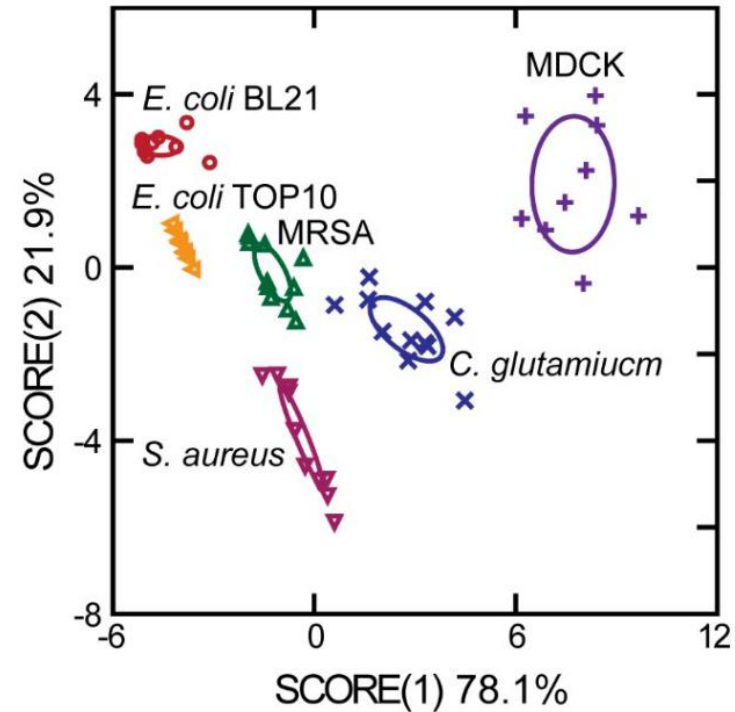


# Selective targeting and detection of bacteria

检测混合菌液中的细菌



**Figure 5b:** Linear discriminant analysis based on the fluorescence response pattern of the 18:1-DNA and 16:0-16:0-DNA conjugates on five types of bacterial strains. The transformed canonical scores were plotted with 95% confidence ellipses around the centroid of each group.



**Figure S11.** Linear discriminant analysis based on the fluorescence response pattern of the 18:1-18:1-DNA and cholesterol-DNA conjugates on five types of bacterial strains and MDCK cells. The transformed canonical scores were plotted with 95% confidence ellipses and 0.001 tolerance around the centroid of each group.