

Miniaturization of CRISPR/Cas12-Based DNA Sensor Array by Non-Contact Printing

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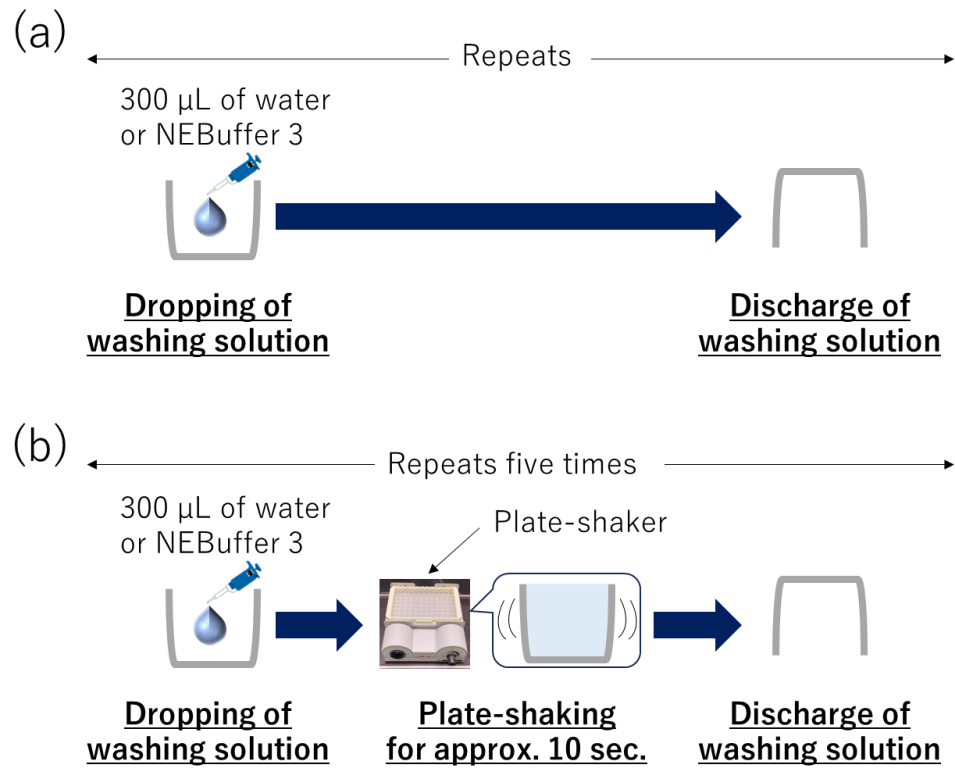


Figure S1. Process to wash the bottom surface of the 96-well: (a) 3.3 section and later, (b) Up to 3.2 section

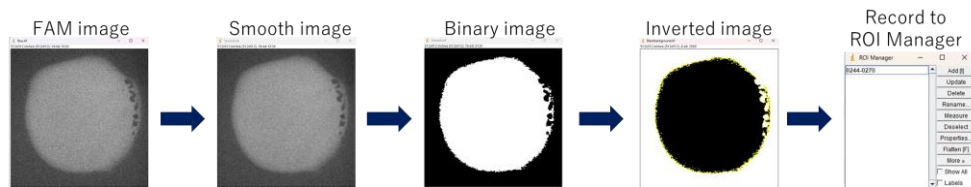


Figure S2. Extraction workflow of the Cas12-immobilized region from the FAM image

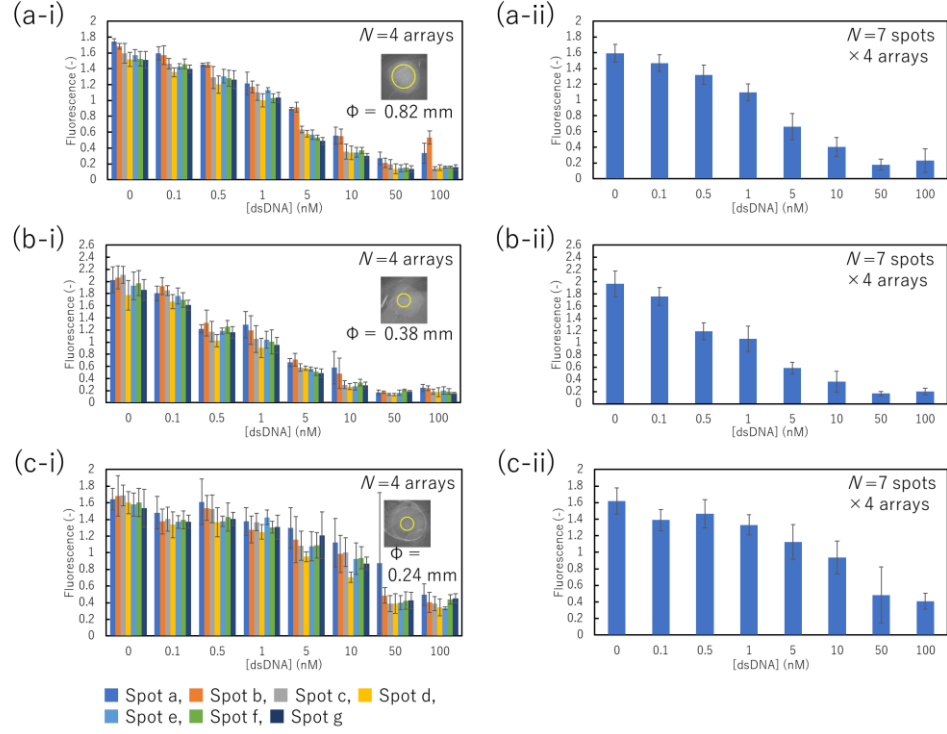


Figure S3. HEX-fluorescence response to target dsDNA concentration in each printing volume condition: (a-i, b-i and c-i) HEX-fluorescence intensity at the center part of each spot (the yellow-circled position) in 40, 20 and 10 nL printing volume condition, respectively, (a-ii, b-ii and c-ii) HEX-fluorescence intensity of each [dsDNA] in 40, 20 and 10 nL printing volume condition, respectively

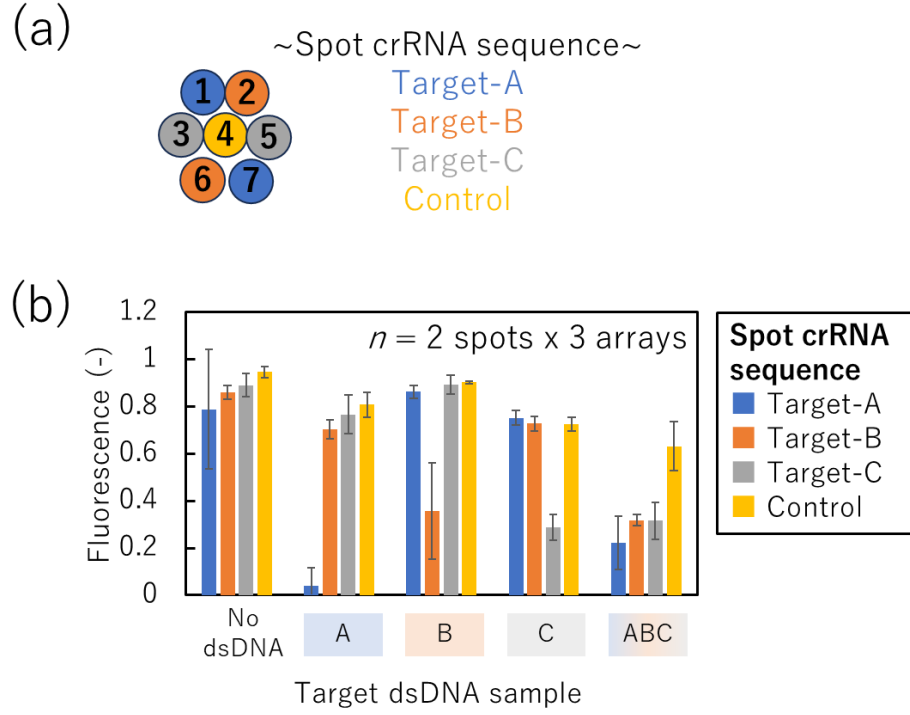


Figure S4. One-pot triple-target dsDNA detection on the non-contact-patterned SPCC-based sensor array fabricated by the optimized process: (a) crRNA sequence of each spot, (b) HEX-fluorescence intensity in each crRNA sequence condition after the incubation of each dsDNA sample (fluorescence intensity was extracted at the region with a radius of 0.71 mm around the center of each spot)

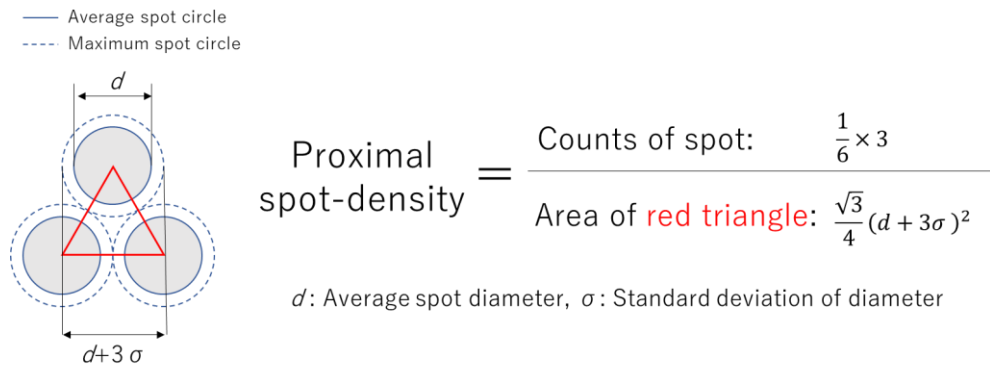


Figure S5. Proximal spot-dispensing model of inkjet patterning and the calculation method of its spot-density

Table S1. Oligonucleotide sequences of amplified target dsDNA, primer, synthesized target dsDNA, crRNA, FAM-crRNA, and ssDNA reporter (the [ACC] sequence of target dsDNA amplicon was converted to [TTT] (part of the PAM sequence) by forward primer; Underline indicates the targeted region of the dsDNA and the target-recognizing region of the crRNA; FAM, HEX, AmC7, and BHQ1 stand for 6-carboxyfluorescein, Hexachloro-fluorescein, Amino Modifier C7 and Black Hole Quencher 1).

Amplified region of target dsDNA (sense strand)	Sequence (5'→3')
pEGFP-N1	CACCTACGGCAAGCTG[ACC]CTGAAGTTCATCTGCACCACCGGCAA-GCTGCCCCGTGCCCTGGCCACCCCTCGTGAC-CACCTGACCTACGGCGTGCACTGCTTCAGCCGCTACCCCGACCACATGAAGCAGCAGCACTTC TTCAAGTCCGCCATGCCCGAAGGCTACGTCCAGGAGCGCACCATCTTCTTCAAGGACG
pVenus-N1	CACCTACGGCAAGCTG[ACC]CTGAAGCTGATCTGCACCACCGGCAA-GCTGCCCCGTGCCCTGGCCACCCCTCGTGAC-CACCTGAGCTACGGCGTGCACTGCTTCGCCCGCTACCCCGACCACATGAAGCAGCAGCACTTC TTCAAGTCCGCCATGCCCGAAGGCTACGTCCAGGAGCGCACCATCTTCTTCAAGGACG
Primer	
Forward	CACCTACGGCAAGCTG[TTT]CTGA
Reverse	CGTCCTTGAAGAAGATGGTGCGC
Synthesized target dsDNA	
Target-A	AATTGCACAATTTGCTCCAAGTGCCTCTGCATTCTTTGGAATGT
Target-B	AGGAGTGTCTTTTCAATTACTTGGGTGTGACCCCTGAAGACTCGGA
Target-C	AATTGCACAATTTGCCCCAGCGCTTCAGCGTTCTTCGGAATGTGCGC
crRNA	
For pVenus-N1	UAAUUUCUACUAAGUGUAGAUUGAAGCUGAUCUGCACC
FAM-crRNA	
For pVenus-N1	UAAUUUCUACUAAGUGUAGAUUGAAGCUGAUCUGCACC -FAM
Negative control	UAAUUUCUACUAAGUGUAGAUUGAAGUAGAUUGGCAGCAC -FAM
For target-A	UAAUUUCUACUAAGUGUAGAUUCCAAGUGCCUCUGCAUUC -FAM
For target-B	UAAUUUCUACUAAGUGUAGAUAAUUACUUGGGUGUGACCCU -FAM
For target-C	UAAUUUCUACUAAGUGUAGAUCCCCAGCGCUUCAGCGUUC -FAM
ssDNA reporter	
HEX-Poly NH2-20nt	T- HEX-TTTTTTTTTTTTTTTTTTTT-AmC7
HEX-Poly NH2-40nt	T- HEX-TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT -AmC7
HEX-Poly NH2-60nt	T- HEX-TT -AmC7
HEX-Poly NH2-80nt	T- HEX-TT -AmC7

HEX-Poly T- HEX-
 NH2-100nt TTTT...TTT-AmC7
 F-Q ssDNA re- HEX-TTATT-BHQ1
 porter

Table S2. Components of PCR mixture. The reagents were added to tubes in order from top to bottom of the list. The 10x Fast Buffer I and dNTP mixture were part of the SpeedSTAR HS Polymerase kit

Reagent	Concentration
Nuclease-free water	—
10x Fast buffer I	1x
dNTP mixture (2.5 mM each)	200 μM
Forward primer	0.4 μM
Reverse primer	0.4 μM
SpeedSTAR HS Polymerase	25 mU μL ⁻¹
pVenus-N1 plasmid	50 fg μL ⁻¹

Table S3. Conditions of non-contact printing of the Cas12-crRNA droplet

Conditions	Before 3.4. section	3.4. and 3.5. section
Syringe pressuring speed	Initiate ($\mu\text{L/s}$)	1 0.2
	Maximum ($\mu\text{L/s}$)	10 1
	Acceleration ($\mu\text{L/s}^2$)	200 100
Pre-pressure before the opening of the solenoid valve (μL)	0.65	
Opening time of the solenoid valve (μs)	650	
Printing volume (nL)	50	10 or 20 or 40

Table S4. Correspondence table of positive (>LOD) / negative (<LOD) samples and their test results, and calculation results of sensitivity (positive results ratio from positive samples) and specificity (negative results ratio from negative samples).

Table S4a. Results in 40 nL-printing volume condition (0, 0.1, 0.5 nM dsDNA samples are defined as negative; 1, 5, 10, 50, 100 nM dsDNA samples are defined as positive)

	Positive samples (>LOD)	Negative Samples (<LOD)
Positive results (>3 σ)	20	1
Negative results (<3 σ)	0	11
	<u>Sensitivity: 100%</u>	<u>Specificity: 92%</u>

Table S4b. Results in 20 nL-printing volume condition (0, 0.1, 0.5 nM dsDNA samples are defined as negative; 1, 5, 10, 50, 100 nM dsDNA samples are defined as positive)

	Positive samples (>LOD)	Negative Samples (<LOD)
Positive results (>3 σ)	20	4
Negative results (<3 σ)	0	8
	<u>Sensitivity: 100%</u>	<u>Specificity: 67%</u>

Table S4c. Results in 10 nL-printing volume condition (0, 0.1, 0.5, 1 nM dsDNA samples are defined as negative; 5, 10, 50, 100 nM dsDNA samples are defined as positive)

	Positive samples (>LOD)	Negative Samples (<LOD)
Positive results (>3 σ)	16	0
Negative results (<3 σ)	0	16
	<u>Sensitivity: 100%</u>	<u>Specificity: 100%</u>