

Supplementary information:

An FcεRI-IgE-based genetically encoded microfluidic cell sensor for fast gram-negative bacterial screening in food samples

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Figures Caption:

Fig. S1. Target sequence design of recombinant protein CD14(Rat)-Fc ϵ IgE.

Fig. S2. (A) Plasmid profile. (B) Agarose gel electrophoresis analysis of CD14(Rat)-IgE Fc plasmid. Lane M: DNA marker; Lane 1: Plasmid of transfection level.

Fig. S3. (A) Identification of protein expression. Lane 1: DMEM of control; Lane 2: DMEM of CD14-Fc ϵ ; Lane M1: SDS-PAGE Marker. (B) SDS-PAGE of affinity purification. Lane M: SDS-PAGE Marker; Lane 1: Supernatant after centrifugation; Lane 2: Effluent after supernatant incubating with Ni-IDA; Lane 3-6: Elution fractions. (C) SDS-PAGE and Western Blot. Left: SDS-PAGE. Right: Western Blot (using anti-His tag antibody). Lane 1: BSA (1.0 μ g); Lane 2: CD14-Fc ϵ protein (1.2 μ g)(Reduced); Lane 3: CD14-Fc ϵ protein (1.2 μ g)(Non-Reduced); Lane M1: SDS-PAGE Marker; Lane M2: Western Blot Marker.

Fig. S4. Standard curve of protein quantification using BCA.

Fig. S5. Immunofluorescence images of CD14-Fc ϵ bound to *E. coli* ATCC 25922. The bacteria were heat-killed and then incubated with (A) or without (B) CD14-Fc ϵ (6 μ g/mL) followed by anti-CD14 (2 μ g/mL) and IgG-FITC.

Fig. S6. Plasmid profile of pLenti-CMV-GCaMP6(s)-2A-Tdtomato

Fig. S7. Lentiviral vector transfection and selection of stable transfection cell lines.

Fig. S8. Cell viability and fluorescence efficiency.

Fig. S9. Fluorescence images of stable transfection RBL-2H3 cell lines after A23187 stimulation.

Fig. S10. β -hexosaminidase assay for confirmation of RBL-2H3 cells activated by *E.coli* ATCC 25922.

Table S1 Comparison of other methods for bacteria detection.

Target protein sequence:

CD14(Rat)-IgE Fc protein Length=604 MW=66008.8 pI=6.78

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1   SPATPEPCEL DQDEESVRCY CNFSDPQPNW SSAFLCAGAE DVEFYGGGRS LEYLLKRVDT
61  EANLGQYTDI IRSLEPLKRLT VRSARVPTQI LFGTLRVLYG SGLRELTLEN LEVTGTALSP
121 LLDATGPDNL TSLLRNVSWA TDTWLAELQ QWLKPKLKV SIAQAHSLNF SCKQVGVFPA
181 LATLDLSDNP ELGKGLISA LCPHKFPTLQ VLALRNAGME TTSGVCSALA AARVPLQALD
241 LSHNSLRDTA GTPSCDWPSQ LNSLNLSFTG LEHVPKGLPA KLSVLDLSYN RLDRKPRPEE
301 LPEVGSLSLT GNPFLHSESQ SEAYNSGVVI ATALSPGSAG LSGTLALLG HRLFVHHHHH
361 HGGGSGGGG SGGGSDDEP RGVITYLIPP SPLDLYENGT PKLTCLVLDL ESEENITVTW
421 VRERKKSIGS ASQRSTKHHN ATTSITSILP VDAKDWIEGE GYQCRVDHPH FPKPIVRSIT
481 KAPGKRSAPE VYVFLPPEEE EKDKRTLCL IQNFFPEDIS VQWLQDSKLI PKSQHSTTPE
541 LKYNQSNQRF FIFSRLEVTK ALWTQTKQFT CRVIHEALRE PRKLERTISK SLGNTSLRPS
601 QASM
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(CD14(Rat) +His +Linker+IgE Fc (CH3+CH4))

Fig. S1. Target sequence design of recombinant protein CD14(Rat)-Fc IgE.

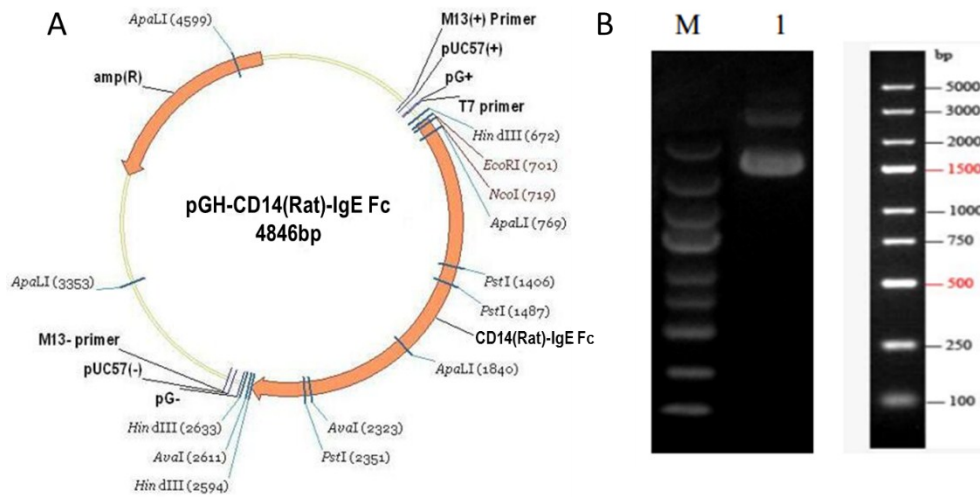


Fig. S2. (A) Plasmid profile. (B) Agarose gel electrophoresis analysis of CD14(Rat)-IgE Fc plasmid. Lane M: DNA marker; Lane 1: Plasmid of transfection level.

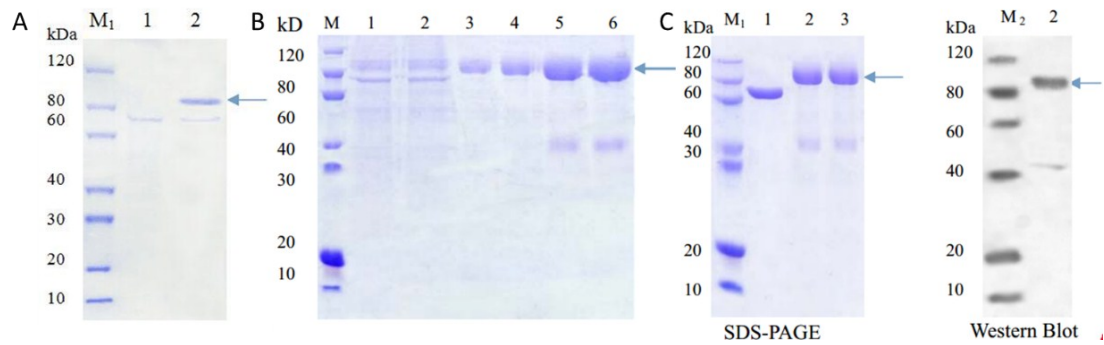


Fig. S3. (A) Identification of protein expression. Lane 1: DMEM of control; Lane 2: DMEM of CD14-Fcε; Lane M1: SDS-PAGE Marker. (B) SDS-PAGE of affinity purification. Lane M: SDS-PAGE Marker; Lane 1: Supernatant after centrifugation; Lane 2: Effluent after supernatant incubating with Ni-IDA; Lane 3-6: Elution fractions. (C) SDS-PAGE and Western Blot. Left: SDS-PAGE. Right: Western Blot (using anti-His tag antibody). Lane 1: BSA (1.0 μg); Lane 2: CD14-Fcε protein (1.2 μg)(Reduced); Lane 3: CD14-Fcε protein (1.2 μg)(Non-Reduced); Lane M1: SDS-PAGE Marker; Lane M2: Western Blot Marker.

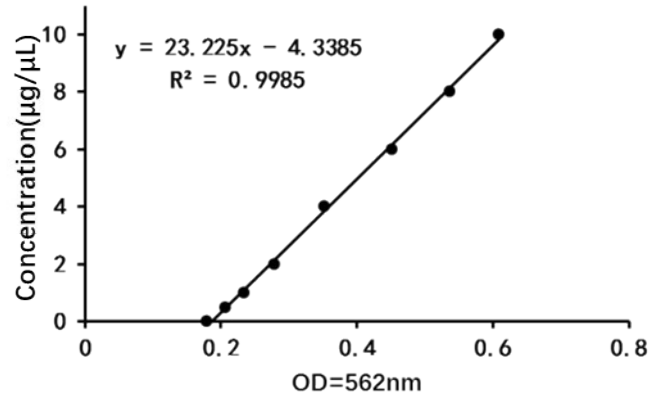


Fig. S4. Standard curve of protein quantification using BCA.

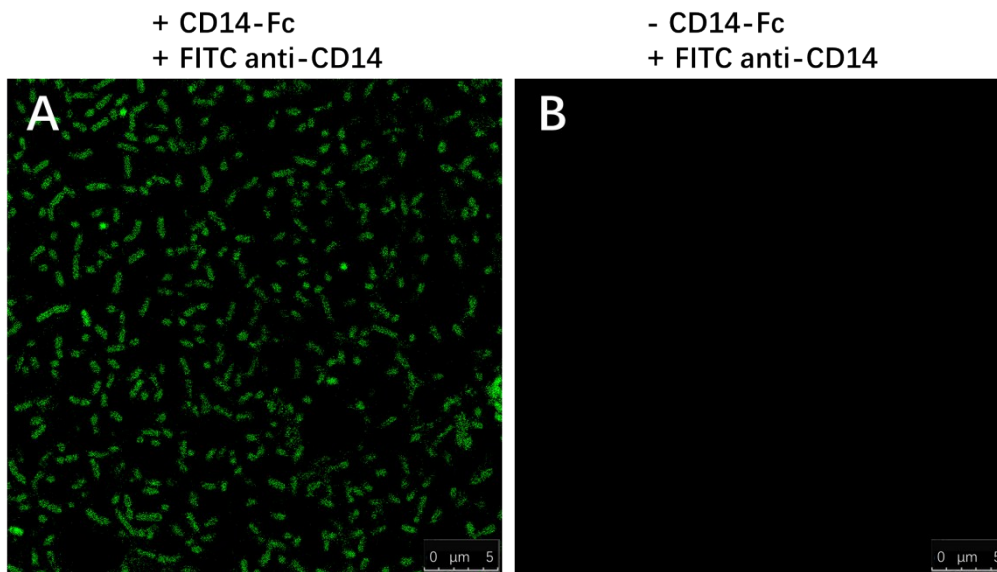


Fig. S5. Immunofluorescence images of CD14-Fc ϵ bound to *E. coli* ATCC 25922. The bacteria were heat-killed and then incubated with (A) or without (B) CD14-Fc ϵ (6 μ g/mL) followed by anti-CD14 (2 μ g/mL) and IgG-FITC.

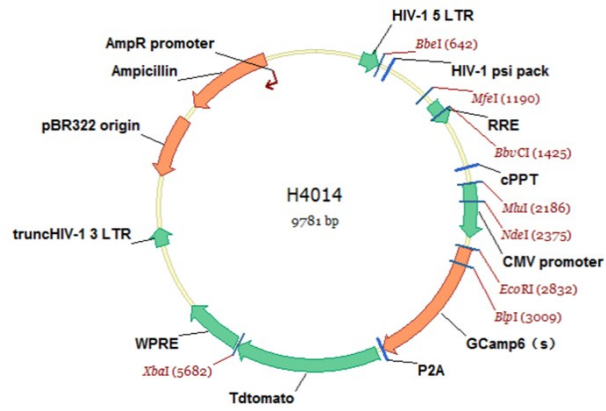


Fig. S6. Plasmid profile of pLenti-CMV-GCaMP6(s)-2A-Tdtomato

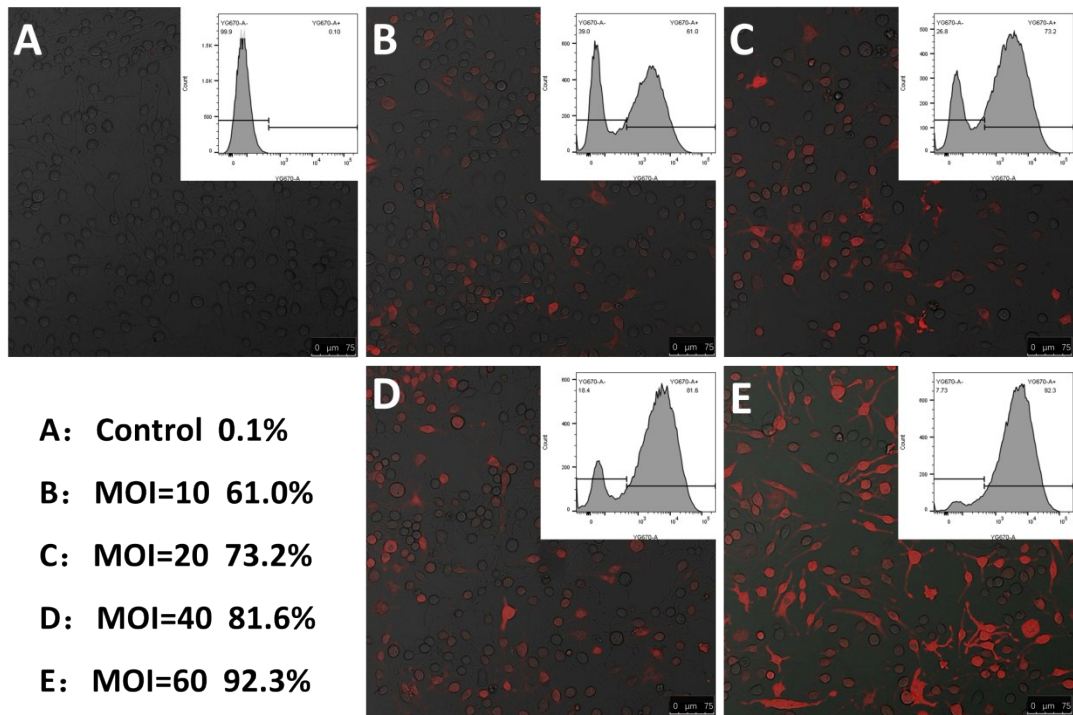


Fig. S7. Lentiviral vector transfection and selection of stable transfection cell lines.

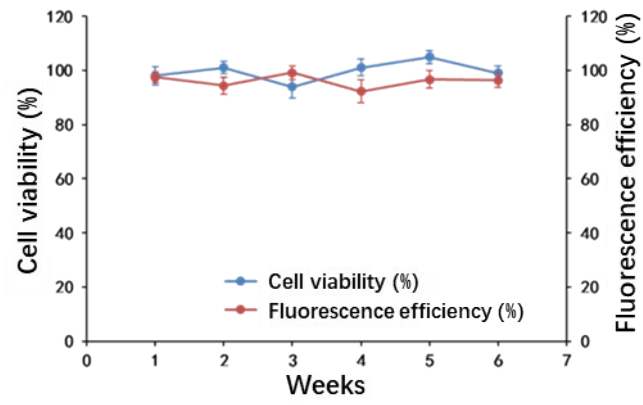


Fig. S8. Cell viability and fluorescence efficiency.

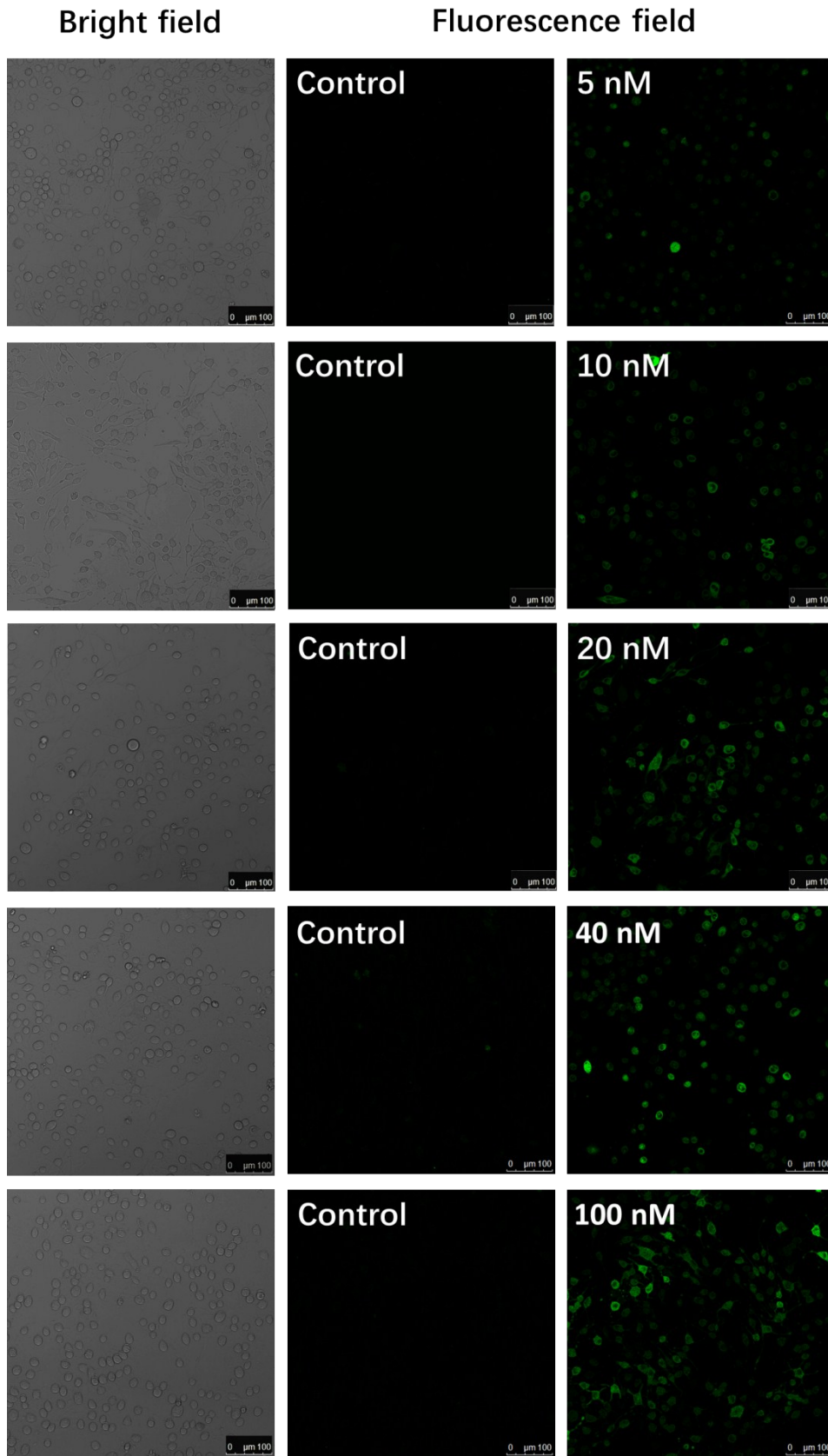


Fig. S9. Fluorescence images of stable transfection RBL-2H3 cell lines after A23187 stimulation.

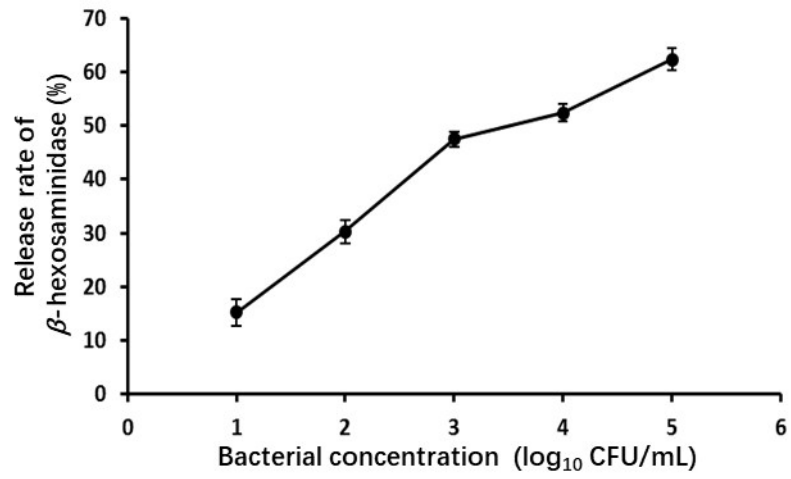


Fig. S10. β -hexosaminidase assay for confirmation of RBL-2H3 cells activated by *E.coli* ATCC 25922.

Table S1 Comparison of other methods for bacteria detection.

Target	Detection method	Linear range (CFU/mL)	LOD (CFU/mL)	Detection time	Ref.
<i>E. coli O157:H7</i>	Impedance based ferrocene-antimicrobial peptide modified biosensor	10^3 – 10^7	10^3	a couple of hours	(Li et al. 2014)
<i>E. coli O157:H7</i>	Aptamer-based QCM sensor	10^2 – 10^7	1.46×10^3	1 h	(Yu et al. 2018)
<i>E. coli O157:H7</i>	Microfluidic colorimetric biosensor	5.0×10^1 – 5.0×10^4	50	45 min	(Zheng et al. 2019)
<i>E. coli O157:H7</i>	Fluorescent biosensor	10 – 10^6	14	2 h	(Xue et al. 2018)
<i>E. coli</i>	Automated microfluidic-based electrochemical sensor	10 – 3.97×10^7	50	8 min	(Altintas et al. 2018)
<i>S. aureus</i>	Aptamer-based hydrogel barcodes	10^2 – 10^4	100	2.5 h	(Xu et al. 2018)
<i>E. faecalis</i>	Molecularly imprinted nanoparticles based plasmonic sensor	2×10^4 – 1×10^8	1.05×10^2	15 min	(Erdem et al. 2019)
<i>S. Enteritidis</i>	Gram staining and direct immunoassay	10^3 – 10^8	800	11 min	(Bu et al. 2019)
<i>V. parahaemolyticus</i>	Aptamer-based visualized detection	10 – 10^6	10	2 h	(Song et al. 2019)
<i>E. coli</i> & <i>S. aureus</i>	Paper-based colorimetric assay	10^4 – 10^8	10^4	20 min	(Sun et al. 2019)
gram-negative bacteria	FcεRI-IgE-based genetically encoded microfluidic cell sensor	8×10^1 – 5×10^3	80	2.5 min	This study

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