

Supplementary Information

A Recombinase-Based Genetic Circuit for Heavy Metal Monitoring

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Table S1. List of oligonucleotides used in this study.

Primer Set	Sequence (5' - 3')	Usage
P_{cadA}-FWD P_{cadA}-REV	TGCCAAGTGAACACCCTGTAGCCAC- TATAGGGTCAAGCCGCCCG- GATGATCCTGAC CTATAGTGGCTACAGGGTGTTTAC- TTGGCAACAGGCGTCGGGGTTT- GTACCGT	P _{cadA} addition to the output module
Bxb1-FWD Bxb1-REV	AGCGGGGTGCACTCATCAAAGAG- GAGAAAGGTACCATGGGAACGGTG TCTTCATGGTAC- CTTTCTCCTCTTTAATACGCGTTTAC- GACATCCCGGTGTG	Overhang primers for Bxb1
CadR-FWD CadR-REV	GAACGGCTACACACCGGGATGTCG- TAAACGCGTATTAAGAGGA- GAAAGGTACCA GAGCCTTTCGTTTATTT- GATGCCCTGCAG- TTAATGCCCCGTGGCTTC	Overhang primers for CadR

Table S2. List of genetic parts and sequences used in this study.

Part	Type	Sequence (5' - 3')
P _{dnaK}	Stress Inducible Promoter	AAAAGCACAAAAAATTTTTGCATCTCCCCCTTGATGACGTGGTTT ACGACCCCATTTAGTAGTCAACCGCAGTGAGTGAGTCTGCAAAA AAATGAAATTGGGCAGTTGAAACCAGACGTTTCGCCCTATTACA GACTCACAACCACATGATGACCGAATATATAGTGAGACGTTTA GATG
mProD	Constitutive Promoter	TCTAGATTTACAGCTAGCTCAGTCCTAGGTATAATGCTAGCTACT AGAG
P _{cadA}	Cadmium Inducible Promoter	TTGACTCTGTAGTTGCTACAGGGTGTGCAAT

IR3	Regulatory	GCAAGCTTGAGCGGGGTGCACTCATCA
Bxb1 attP	Regulatory	GTCGTGGTTTGTCTGGTCAACCACCGCGGTCTCAGTGGTGTACGG TACAAACCCCGAC
Bxb1 attB	Regulatory	GCCCCGATGATCCTGACGACGGAGACCGCCGTCGTCGACAAGCC GGCCGA
RBS	Regulatory	AAAGAGGAGAAA
rrnBT1	Terminator	CAAATAAAACGAAAGGCTCAGTCGAAAGACTGGGCCTTTCGTTT TATCTGTTGTTTGTCCGGTGAACGCTCTCCTGAGTAGGACAAAT ATGCGTAAAGGAGAAGAAGCTTTTCACTGGAGTTGTCCCAATTCTT GTTGAATTAGATGGTGTATGTTAATGGGCACAAATTTTCTGTCTCAGT GGAGAGGGTGAAGGTGATGCAACATACGGAAAACCTTACCCTTAA ATTTATTTGCACTACTGGAAAACCTGTTCCATGGCCAACACTT GTCACTACTTTTCGGTTATGGTGTTCATGCTTTGCGAGATACCCAG ATCATATGAAACAGCATGACTTTTTCAAGAGTGCCATGCCCGAAG GTTATGTACAGGAAAGAACTATATTTTTCAAAGATGACGGGAACT ACAAGACACGTGCTGAAGTCAAGTTTGAAGGTGATACCCTTGTTA ATAGAATCGAGTTAAAAGGTATTGATTTTAAAGAAGATGGAAAC ATTCTTGACACAAATTGGAATACAACATAACTCACACAATGTA TACATCATGGCAGACAAACAAAAGAATGGAATCAAAGTTAACTT CAAAATTAGACACAACATTGAAGATGGAAGCGTTCAACTAGCAG ACCATTATCAACAAAATACTCCAATTGGCGATGGCCCTGTCTTT TACCAGACAACCATTACCTGTCCACACAATCTGCCCTTTCGAAAG ATCCCAACGAAAAGAGAGACCACATGGTCTTCTTGAGTTTGTA CAGCTGCTGGGATTACACATGGCATGGATGAACTATACAAATAA ATGGCCAAAAATCCGAAAGATGGCGAAAGCCGCACCTTCCTGAT TAGCGTGGCCGCCGAAGTGGCCGGTATGCATGCCAGACCCTGC GCACCTATGATCGTCTGGGTCTGGTGAGCCCGCGTCGTACCAGTG GTGGTGGTCGTCTGTTATAGCCTGCATGATGTGGAGCTGCTGCGCC AGGTTACAGCATCTGAGCCAGGATGAAGGCGTGAATCTGGCCGGC ATCAAACGCATCATTGAACTGACCAGCCAGGTGGAAGCACTGCA GAGCCGCCTGCAGGAAATGGCCGAAGAACTGGCCGTGCTGCGCG CCAATCAGCGTCGTGAAGTGGCCGTGGTGCCGAAAAGCACCGCC CTGGTGGTGTGGAACCGCGTCGTTAA
GFP	CDS	ATGAAGATCGGAGAACTGGCCAAAGCCACCGACTGCGCCGTGGA AACCATCCGCTACTACGAGCGTGAACAGCTGCTGCCGGAGCCGG CACGCAGCGACGGCAACTACCGGCTGTACACCCAGGCCACGTC GAGCGGCTTACCTTCATCCGCAACTGCCGCACCCTGGACATGACC CTGGATGAAATCCGCAGCCTGCTACGCCTGCGCGACAGCCCCGA TGATTTCGTGCGGCAGCGTCAATGCGCTGATCGACGAGCATATCGA GCATGTGCAGGCACGGATCGATGGTCTGGTGGCGTTGCAGGAAC AGCTGGTGGAGCTGCGGCGGCGCTGCAATGCACAAGGGGCGGA GTGTGCGATCTTGCAGCAACTGGAGACGAACGGGGCGGTATCGG TGCCGGAACCGAGCATTGCGATGTAGGGCGAAGCCACGGGCAT TAA
HspR	CDS	ATGGGAACGGTGGCGCAGATGGAATTAGAAGCGATCAAAGAGC GGAACCGTTCGGCTGCGCATTTCATATCCGCGCCGGGAAATACC GAGGATCCCTGCCGCCGTGGGGATACCTGCCTACGCGCGTGGAC GGGGAGTGGCGGCTGGTGGCCGACCTGTGCAGCGAGAGCGCAT CCTCGAGGTGTATCACCGCGTCGTCGACAACCACGAGCCGCTGC ATCTGGTGGCCACGACCTGAACCGGCGTGGTGTCTGTGCGCGA AGGACTACTTCGCGCAGCTGCAAGGCCGCGAGCCGAGGGCCGG
CadR	CDS	ATGGGAACGGTGGCGCAGATGGAATTAGAAGCGATCAAAGAGC GGAACCGTTCGGCTGCGCATTTCATATCCGCGCCGGGAAATACC GAGGATCCCTGCCGCCGTGGGGATACCTGCCTACGCGCGTGGAC GGGGAGTGGCGGCTGGTGGCCGACCTGTGCAGCGAGAGCGCAT CCTCGAGGTGTATCACCGCGTCGTCGACAACCACGAGCCGCTGC ATCTGGTGGCCACGACCTGAACCGGCGTGGTGTCTGTGCGCGA AGGACTACTTCGCGCAGCTGCAAGGCCGCGAGCCGAGGGCCGG
Bxb1	CDS	ATGGGAACGGTGGCGCAGATGGAATTAGAAGCGATCAAAGAGC GGAACCGTTCGGCTGCGCATTTCATATCCGCGCCGGGAAATACC GAGGATCCCTGCCGCCGTGGGGATACCTGCCTACGCGCGTGGAC GGGGAGTGGCGGCTGGTGGCCGACCTGTGCAGCGAGAGCGCAT CCTCGAGGTGTATCACCGCGTCGTCGACAACCACGAGCCGCTGC ATCTGGTGGCCACGACCTGAACCGGCGTGGTGTCTGTGCGCGA AGGACTACTTCGCGCAGCTGCAAGGCCGCGAGCCGAGGGCCGG

GAGTGGTCGGCTACCGCGCTGAAGCGATCGATGATCTCCGAGGC
 GATGCTCGGGTACGCGACTCTGAACGTAAGACCGTCCGAGACG
 ACGACGGAGCCCCGCTGGTGCGGGCTGAGCCGATCCTGACCCGT
 GAGCAGCTGGAGGCGCTGCGCGCCGAGCTCGTGAAGACCTCCCC
 GGCGAAGCCCCGCGGTGTCTACCCCGTCGCTGCTGCTGCGGGTGT
 GTTCTGCGCGGTGTGCGGGGAGCCCGCGTACAAGTTCGCCGGGG
 GAGGACGTAAGCACCCGCGCTACCGCTGCCGCTCGATGGGGTTC
 CCGAAGCACTGCGGGAACGGCACGGTGGCGATGGCCGAGTGGG
 ACGCGTTCTGCGAGGAGCAGGTACTGGATCTGCTCGGGGACGCG
 GAGCGTCTGGAGAAAGTCTGGGTAGCGGGCTCGGACTCCGCGGT
 CGAACTCGCGGAGGTGAACGCGGAGCTGGTGGACCTGACGTCGC
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 GCACTGGATGCCCCGTATTGCGGCGCTGGCCGCGCGGCAAGAGGA
 GCTGGAGGGCCTGGAGGCTCGCCCGTCTGGCTGGGAGTGGCGCG
 AGACCGGGCAGCGGTTCCGGGACTGGTGGCGGGAGCAGGACAC
 CGCGGCAAAGAACACCTGGCTTCGGTCGATGAACGTTCCGGCTGA
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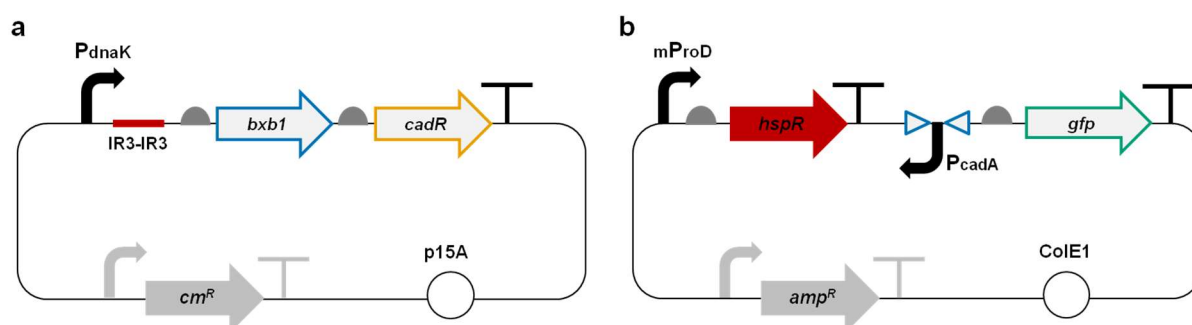


Figure S1. Representative plasmid maps of the cadmium sensor. Plasmid map containing actuating module (a) and sensing and output modules (b). “T” represents terminator.

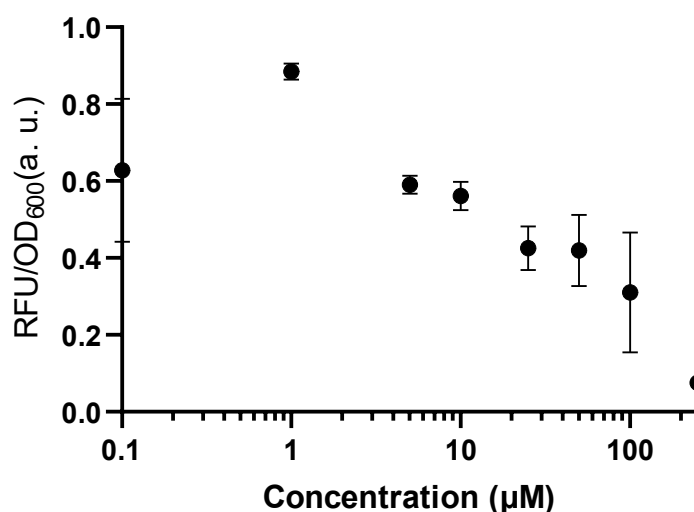


Figure S2. Cadmium-related toxicity for Escherichia coli DH5α strain. The initial number of cells was 1:50 of culture volume. The cells were cultured in MOPS minimal media with a pH of 7.5 at 37°C for 14h. Measurements were done on 96-well plates. The fluorescence intensity of each group

was compared with each other and normalized according to the formula stated in the Materials and Methods section. Data were visualized with mean \pm standard error mean (SEM) in each graph. At least three biological replicates were used for each analysis.

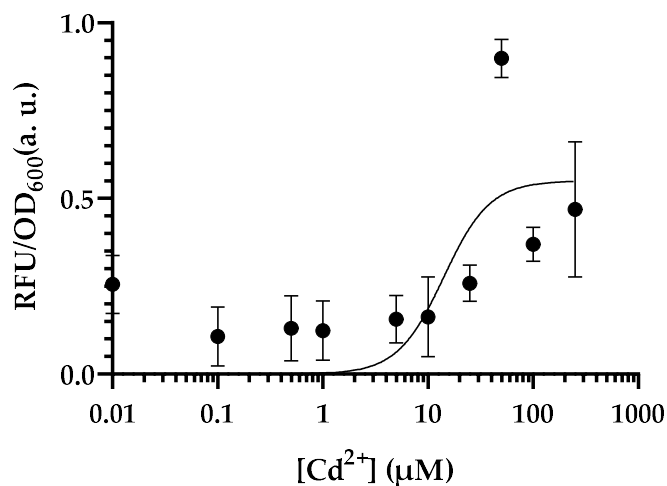


Figure S3. The system response with 100 μM and 250 μM . The initial number of cells was 1:50 of culture volume. The cells were cultured in MOPS minimal media with a pH of 7.5 at 37°C for 10h. Measurements were done on 96-well plates. The fluorescence intensity of each group was compared with each other and normalized according to the formula stated in the Materials and Methods section. Data were visualized with mean \pm standard error mean (SEM) in each graph. At least three biological replicates were used for each analysis.