

# Visualizing the Perturbation of Cellular Cyclic di-GMP Levels in Bacterial Cells

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Supporting Information

ABSTRACT: Cyclic di-GMP (c-di-GMP) has emerged as a prominent intracellular messenger that coordinates biofilm formation and pathogenicity in many bacterial species. Developing genetically encoded biosensors for cdi-GMP will help us understand how bacterial cells respond to environmental changes via the modulation of cellular c-di-GMP levels. Here we report the design of two genetically encoded c-di-GMP fluorescent biosensors with complementary dynamic ranges. By using the biosensors, we found that several compounds known to promote biofilm dispersal trigger a decline in c-di-GMP levels in Escherichia coli cells. In contrast, cellular c-di-GMP levels were elevated when the bacterial cells were treated with subinhibitory concentrations of biofilm-promoting antibiotics. The biosensors also revealed that E. coli cells engulfed by macrophages exhibit lower c-di-GMP levels, most likely as a response to the enormous pressures of survival during phagocytosis.

In recent years, the cyclic dinucleotide c-di-GMP has emerged as a prominent messenger that coordinates the cellular functions associated with bacterial biofilm formation and pathogenicity.<sup>1-3</sup> In bacterial cells, the concentration of c-di-GMP is controlled by a large number of diguanylate cyclases (DGCs) and c-di-GMP phosphodiesterases (PDEs) (Figure 1a). C-di-GMP exerts its effect by binding to a diverse array of receptors, including enzymes, transcriptional factors, adaptor proteins, and riboswitches.<sup>4</sup> It has become increasingly clear that c-di-GMP signaling networks play central roles in bacterial biofilm formation and virulence gene expression in some clinically important pathogens, such as *Vibrio cholerae*, *Pseudomonas aeruginosa*, and *Klebsiella pneumoniae*.<sup>5-8</sup>

One of the most remarkable features of c-di-GMP signaling networks is that they usually consist of a large number of DGCs and PDEs, with many of them containing putative sensory domains for perceiving environmental cues.<sup>9–12</sup> It is believed that the diverse array of sensory domains allows the bacterial cells to respond to environmental changes by modulating the cellular c-di-GMP concentration through the DGC and PDE proteins. However, with a few exceptions, the vast majority of the environmental signals and associated c-di-GMP pathways remain to be unveiled. In this regard, developing biosensors that can report the changes in cellular c-di-GMP levels would facilitate the identification of the proteins and pathways involved in the



**Figure 1.** Design and in vitro characterization of c-di-GMP biosensors. (a) Synthesis and degradation of c-di-GMP by DGCs and PDEs. (b) Construction of the genetically encoded FRET-based biosensors for c-di-GMP using MrkH and VCA0042. Both proteins contain a c-di-GMP binding PilZ domain and an N-terminal domain (NTD). (c, d) Fluorescence titration curves for cdg-S1 and cdg-S2 (experimental conditions are included in the SI). (e) Schematic illustration of the conformational change induced by binding c-di-GMP to cdg-S1 and cdg-S2.

response mechanisms that are essential for environmental adaptation.

We previously reported a fluorescent-dye-labeled c-di-GMP biosensor for in vitro c-di-GMP detection that uses the nonenzymatic EAL domain of the FimX protein.<sup>13,14</sup> Here we describe the design of two genetically encoded fluorescent biosensors for monitoring cellular c-di-GMP concentrations using two natural c-di-GMP binding proteins. The Förster resonance energy transfer (FRET)-based fluorescent biosensors were constructed by using MrkH and VCA0042, two c-di-GMP binding proteins from *K. pneumoniae* and *V. cholerae*,

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respectively.<sup>6,15</sup> The two proteins share a common C-terminal PilZ domain for c-di-GMP binding but contain different Nterminal domains (NTDs). The genes encoding the two proteins were cloned into pET28b- and pUCP18-based expression vectors that harbor the mCerulean and mVenus genes to produce cdg-S1 and cdg-S2 (Figure 1b) after optimization of the length of the linkers flanking the mrkH and VCA0042 genes. mCerulean and mVenus were derived from the standard cyan fluorescent protein (CFP) and yellow fluorescent protein (YFP) to generate more stable fluorescent proteins with brighter fluorescence.<sup>16</sup> For in vitro characterization of the biosensors, cdg-S1 and cdg-S2 were produced using the Escherichia coli expression system and treated with the protein RocR to remove the associated c-di-GMP.<sup>17</sup> Upon full maturation of mCerulean and mVenus, fluorescence titrations were performed to show that the addition of c-di-GMP gradually reduces the relative change in emission ratio (Figure 1c,d), which implies a reduction in FRET efficiency. On the basis of the crystal structure of VCA0042,<sup>15</sup> the reduced FRET efficiency is likely caused by a ligand-induced conformational change that affects the dipoledipole orientation (or distance) between mCerulean and mVenus (Figure 1e). The reduction in FRET efficiency upon c-di-GMP binding is also reminiscent of that observed with the biosensor developed from the protein YcgR.<sup>18</sup> Fitting of the titration data yielded dissociation constants  $(K_d)$  of 0.12 and 2.4  $\mu$ M for cdg-S1 and cdg-S2, respectively. In view of the fact that the estimated concentrations of c-di-GMP in bacterial cells are in the range  $0.1-10 \,\mu$ M, the complementary dynamic ranges of the two biosensors should allow the reporting of cellular c-di-GMP levels at both the low and high ends of the concentration gradient. The presence of other nucleotides (e.g., cGMP, GXP, AXP, NADP<sup>+</sup>, etc.) at physiologically relevant concentrations did not interfere with the performance of the biosensors [see Table S1 in the Supporting Information (SI)]. The two biosensors are also insensitive to pH changes in the pH range from 6 to 9 (data not shown).

When cdg-S1 and cdg-S2 were expressed in the BL21 *E. coli* strain by isopropylthio- $\beta$ -galactoside (IPTG) induction or constitutively in the uropathogenic *E. coli* strain UTI89, cyan and yellow fluorescence could be readily detected using a confocal microscope. Interestingly, in contrast to a diffuse distribution of the two biosensors in the *E. coli* UTI89 cells, most of the biosensors appeared to be sequestered to the poles of the BL21 cells regardless of expression level. We found that the single-expressed mVenus or mCerulean also has a strong tendency to cluster at the poles in BL21 cells (Figure S1 in the SI), indicating that the polar localization could be due to the interaction between mVenus and unknown *E. coli* polar proteins. As we demonstrate below, similar results were obtained from the two *E. coli* strains regardless of the polarization of the biosensors.

A group of structurally diverse compounds are known to hinder the formation of robust bacterial biofilm or trigger the dispersal of biofilm.<sup>19–24</sup> It is also known that biofilm dispersal can be induced by sequestering cellular c-di-GMP by overexpressing c-di-GMP binding proteins, which indicates that a low c-di-GMP level could directly lead to biofilm dispersal.<sup>25</sup> By using the biosensors, we asked whether some of the biofilm-dispersing compounds (or dispersal factors) can cause a reduction in cellular c-di-GMP levels in planktonic *E. coli* cells. The answer to the question would yield insight into the biofilm- dispersing mechanism of these compounds. The compounds under investigation included one of the *N*-acylhomoserine lactone (AHL) autoinducers (*N*-hexanoyl-DL-homoserine lactone), the plant auxin 3-indolyacetonitrile (3-IAN),<sup>23</sup> D-tyrosine,<sup>20</sup> resveratrol,<sup>26</sup> and the nitric oxide (NO) donor MAHMA-NONOate.<sup>21</sup> As shown in Figure 2, Figure S2, and Table S2, an increase in



**Figure 2.** Perturbation of c-di-GMP levels in BL21 *E. coli* cells by biofilm-dispersing agents. (a) Average FRET efficiencies for the *E. coli*-containing biosensors before and after the treatment with biofilm-dispersing agents for 30 min. The FRET efficiency was measured by observing the change in donor emission (466 nm) upon acceptor photobleaching (see the SI). Statistical significance is indicated by the asterisks (\*, *P* < 0.05; \*\*, *P* < 0.01). Changes in FRET efficiency are directly correlated to changes in the population of ligand binding biosensors and thus to the c-di-GMP level. (b) Time-dependent changes in the FRET efficiencies for 3-IAN-treated cells. (c) Representative images of YFP ( $\lambda_{ex} = 430$  nm,  $\lambda_{em} = 528$  nm) prior to acceptor photobleaching.

FRET efficiency was observed for four of the five compounds in both E. coli strains. D-Tyrosine was the only compound that did not seem to perturb the FRET efficiency. A time-dependent study of 3-IAN showed an overall increase in FRET efficiency over time, indicating that the c-di-GMP level decreased gradually after an initial increase (Figure 2b,c and Figure S2b,c). In comparison, the control experiment showed that the FRET efficiency did not change significantly for either E. coli strain during the 1 h observation window in the absence of dispersal factors (Figure 2c). It should be noted that cdg-S1 and cdg-S2 exhibited similar response dynamics but different dynamic ranges (Figure 2b), indicating that the cellular c-di-GMP concentration is likely in the range 0.1–10  $\mu$ M. On the basis of the standard curves obtained from in vitro titration data, we estimated that the compounds caused a drop of cellular c-di-GMP concentration from 0.8–2  $\mu$ M to 0.2–0.5  $\mu$ M. These results indicate that some of the biofilm-dispersing compounds may indeed induce biofilm dispersal by reducing cellular c-di-GMP concentration. The observation that D-tyrosine did not perturb the c-di-GMP level is in accordance with the view that D-amino acids trigger biofilm disassembly through the replacement of D-alanine in the biosynthesis of cell-wall fibers, a process that does not involve the cytoplasmic c-di-GMP messenger.<sup>20</sup> The effect of NO on the c-di-GMP level is probably exerted through the direct regulation of the DGC or PDE proteins.<sup>21,27</sup> The detailed molecular mechanisms for the other dispersal factors remain to be determined.

In contrast to the dispersal factors, studies have shown that subinhibitory concentrations of aminoglycosides and cell-wall-targeting antibiotics can promote biofilm formation.<sup>28–31</sup> In view of the central role of c-di-GMP in biofilm formation, it was speculated that the biofilm-promoting effect of some of the antibiotics is exerted through c-di-GMP. The biosensors allowed us to test directly whether treatment with subinhibitory concentrations of antibiotics could raise cellular c-di-GMP concentrations. As shown in Figure 3, Figure S3, and Table S3,



**Figure 3.** Perturbation of c-di-GMP levels in BL21 *E. coli* cells by antibiotics at subinhibitory concentrations. (a) Average FRET efficiencies for the *E. coli*-containing biosensors before and after treatment with the antibiotics for 30 min. The FRET efficiency was measured by observing the change in donor emission (466 nm) upon acceptor photobleaching (see the SI). Statistical significance is indicated by the asterisks (\*, p < 0.05; \*\*, P < 0.01). Data were obtained by averaging the readings for multiple cells (n > 10). Changes in the FRET ratio are directly correlated to changes in the population of ligand-binding biosensors and thus to the c-di-GMP level. (b) Time-dependent changes in the FRET efficiencies for gentamycin. (c) Representative images of YFP ( $\lambda_{ex} = 430$  nm,  $\lambda_{em} = 528$  nm) prior to acceptor photobleaching.

when the E. coli cells were treated with the representative antibiotics at subinhibitory concentrations, significant decreases in FRET efficiency were observed for some of the antibiotics. Overall, the aminoglycoside antibiotics (tobramycin, gentamicin, and streptomycin) and macrolide antibiotic (erythromycin) that target ribosome and the antibiotics (ampicillin and vancomycin) that target cell-wall biosynthesis caused substantial reductions in the FRET efficiency, indicating an increase in cellular c-di-GMP concentration. In contrast, the two antibiotics that are not known to induce biofilm formation (mitomycin C and norfloxacin) did not seem to change the c-di-GMP level. These observations provide support for the view that subinhibitory concentrations of antibiotics trigger biofilm formation by raising the cellular c-di-GMP concentration. Notably, the role of c-di-GMP in aminoglycoside-induced biofilm formation was unveiled previously by the discovery that aminoglycosides induce the upregulation of YdeH, a DGC that produces c-di-GMP in E.

*coli.*<sup>30</sup> A c-di-GMP-specific PDE from *P. aeruginosa* has also been found to play a crucial role in tobramycin-inducible biofilm formation.<sup>29</sup>

The time-dependent change in the FRET efficiency during gentamycin treatment revealed the dynamics of the response to the antibiotic. The results showed a rapid rise in the c-di-GMP level in the first 10 min of treatment followed by a gradual decrease (Figure 3b and Figure S3b). It should be noted that the antibiotic treatment reduced the FRET signal of cdg-S1 to very low levels and that the recovery of the FRET signal was consistently much slower than for cdg-S2. In the UT189 strain, the difference between the two biosensors was not as prominent (Figure S3b). The discrepancies between the two biosensors in the two E. coli strains are likely caused by the different dynamic ranges of the biosensors and higher c-di-GMP levels in the UT189 cells under antibiotic stresses. On the basis of these results, we estimated that the biofilm-promoting antibiotics can raise the cellular c-di-GMP concentration in BL21 cells to as high as 5–10  $\mu$ M from the basal level of 0.8–2  $\mu$ M. The c-di-GMP levels in the UT189 cells could be even higher, with both the cdg-S1 and cdg-S2 saturated upon antibiotic treatment.

Lastly, c-di-GMP has been suggested to play a role in the in vivo survival of some intracellular pathogenic bacteria.<sup>32,33</sup> With the biosensor, we investigated whether the c-di-GMP level changed when bacterial cells were challenged by the host immune system. One of the most challenging environments for bacterial cells is inside the phagosomes of macrophages during the phagocytosis process. During phagocytosis, the bacterial cells are under enormous stress in a low-pH environment that is inundated with a host of antimicrobial agents such as NO, reactive oxygen species (ROS), and antimicrobial peptides. By averaging the readings from multiple bacterial cells, we observed increased FRET efficiencies for both cdg-S1 and cdg-S2 when the E. coli cells were engulfed by RAW 264.7 macrophage cells (Figure 4 and Figure S4). The increase in FRET efficiency upon engulfment can also be seen from the histogram of the free and engulfed cells (Figure 4c and Figure S4c; only the cdg-S2 results are shown). These observations indicate a reduction in c-di-GMP concentration in the engulfed E. coli cells, similar to the effect triggered by the biofilm-dispersing compounds (e.g., NO) discussed above. The reduced c-di-GMP level in the macrophage-engulfed cells is most likely part of the counterattack strategy used by the bacterial cells to boost virulence expression to cope with the enormous pressure for survival. This view is further supported by the observation that a decrease in c-di-GMP level in the intracellular pathogen Bordetella is correlated with its ability to kill macrophages.<sup>34</sup>

In summary, we have developed two genetically encoded FRET biosensors for monitoring the fluctuation of c-di-GMP levels in bacterial cells. The biosensors revealed a decline of c-di-GMP level when the cells were treated with biofilm-dispersing agents or in the hostile environment of macrophages. On the contrary, the biosensors reported elevated c-di-GMP levels in E. coli when the cells were treated with subinhibitory concentrations of biofilm-promoting antibiotics. These observations support the view that high c-di-GMP levels promote sessility and biofilm formation while low c-di-GMP levels promote biofilm dispersal. The results establish these biosensors as valuable tools for use in chemical biology and indicate the crucial role played by c-di-GMP in stress response and environmental adaptation. In conjunction with studies on mutant strains, the biosensors will in the future further help us to identify the specific c-di-GMP signaling proteins and pathways involved in stress response. By

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**Figure 4.** Perturbation of c-di-GMP levels in BL21 *E. coli* cells engulfed by macrophages. (a) Representative images of YFP ( $\lambda_{ex} = 430 \text{ nm}$ ,  $\lambda_{em} = 528 \text{ nm}$ ) prior to acceptor photobleaching of the *E. coli* cells outside and inside RAW264.7 macrophage cells. The side panels show enlarged views of the bacterial cells. (b) Average FRET efficiencies for *E. coli* cells outside and inside the macrophage cells. Statistical significance is indicated by the asterisks (\*\*\*, *P* < 0.001). (c) Histogram showing the FRET efficiency distribution for cdgS2 in the free and macrophage engulfed *E. coli* cells. The total numbers of cells were 50 (free) and 53 (engulfed).

studying the biosensor-containing bacterial cells embedded in biofilm matrixes, we will also be able to gain a better understanding of the roles of c-di-GMP in the highly dynamic and complex processes of biofilm formation and dispersal.

# ASSOCIATED CONTENT

### **Supporting Information**

Experimental details and supporting figures and tables. This material is available free of charge via the Internet at http://pubs. acs.org.

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### Notes

The authors declare no competing financial interest.

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#### REFERENCES

(1) Hengge, R. Nat. Rev. Microbiol. 2009, 7, 263.

- (2) Jenal, U.; Malone, J. Annu. Rev. Genet. 2006, 40, 385.
- (3) Romling, U.; Amikam, D. Curr. Opin. Microbiol. 2006, 9, 218.

(4) Ryan, R. P.; Tolker-Nielsen, T.; Dow, J. M. Trends Microbiol. 2012, 20, 235.

(5) Kulasakara, H.; Lee, V.; Brencic, A.; Liberati, N.; Urbach, J.; Miyata, S.; Lee, D. G.; Neely, A. N.; Hyodo, M.; Hayakawa, Y.; Ausubel, F. M.; Lory, S. *Proc. Natl. Acad. Sci. U.S.A.* **2006**, *103*, 2839.

(6) Wilksch, J. J.; Yang, J.; Clements, A.; Gabbe, J. L.; Short, K. R.; Cao, H.; Cavaliere, R.; James, C. E.; Whitchurch, C. B.; Schembri, M. A.; Chuah, M. L. C.; Liang, Z.-X.; Wijburg, O. L.; Jenney, A. W.; Lithgow, T.; Strugnell, R. A. *PLoS Pathog.* **2011**, *7*, No. e1002204.

(7) Tischler, A. D.; Camilli, A. Infect. Immun. 2005, 73, 5873.

(8) Lim, B.; Beyhan, S.; Meir, J.; Yildiz, F. H. Mol. Microbiol. 2006, 60, 331.

(9) Galperin, M. Y.; Nikolskaya, A. N.; Koonin, E. V. FEMS Microbiol. Lett. 2001, 203, 11.

(10) Qi, Y.; Rao, F.; Liang, Z.-X. Biochemistry 2009, 48, 10275.

(11) Tarutina, M.; Ryjenkov, D. A.; Gomelsky, M. J. Biol. Chem. 2006, 281, 34751.

(12) Delgado-Nixon, V. M.; Gonzalez, G.; Gilles-Gonzalez, M. A. Biochemistry 2000, 39, 2685.

(13) Ho, C. L.; Koh, S. L.; Chuah, M. L. C.; Luo, Z.; Tan, W. J.; Low, D. K. S.; Liang, Z.-X. *ChemBioChem* **2011**, *12*, 2753.

(14) Qi, Y.; Chuah, M. L. C.; Dong, X.; Xie, K.; Luo, Z.; Tang, K.; Liang, Z.-X. J. Biol. Chem. **2011**, 286, 2910.

(15) Benach, J. EMBO J. 2007, 26, 5153.

(16) Markwardt, M. L.; Kremers, G. J.; Kraft, C. A.; Ray, K.; Cranfill, P. J. C.; Wilson, K. A.; Day, R. N.; Wachter, R. M.; Davidson, M. W.; Rizzo, M. A. *PLoS One* **2011**, *6*, No. e17896.

(17) Chen, M. W.; Kotaka, M.; Vonrhein, C.; Bricogne, G.; Rao, F.; Chuah, M. L. C.; Svergun, D.; Schneider, G.; Liang, Z.-X.; Lescar, J. J. Bacteriol. **2012**, 194, 4837.

(18) Christen, M.; Kulasekara, H. D.; Christen, B.; Kulasekara, B. R.; Hoffman, L. R.; Miller, S. I. *Science* **2010**, 328, 1295.

(19) McDougald, D.; Rice, S. A.; Barraud, N.; Steinberg, P. D.; Kjelleberg, S. Nat. Rev. Microbiol. 2012, 10, 39.

(20) Kolodkin-Gal, I.; Romero, D.; Cao, S.; Clardy, J.; Kolter, R.; Losick, R. *Science* **2010**, 328, 627.

(21) Barraud, N.; Schleheck, D.; Klebensberger, J.; Webb, J. S.; Hassett, D. J.; Rice, S. A.; Kjelleberg, S. *J. Bacteriol.* **2009**, *191*, 7333.

(22) Kolodkin-Gal, I.; Čao, S.; Chai, L.; Böttcher, T.; Kolter, R.; Clardy, J.; Losick, R. *Cell* **2012**, *149*, 684.

(23) Frei, R.; Breitbach, A. S.; Blackwell, H. E. Angew. Chem., Int. Ed. 2012, 51, 5226.

(24) Waters, C. M.; Lu, W.; Rabinowitz, J. D.; Bassler, B. L. J. Bacteriol. 2008, 190, 2527.

(25) Ma, Q.; Yang, Z.; Pu, M.; Peti, W.; Wood, T. K. *Environ. Microbiol.* **2011**, *13*, 631.

(26) Coenye, T.; Brackman, G.; Rigole, P.; De Witte, E.; Honraet, K.; Rossel, B.; Nelis, H. J. *Phytomedicine* **2012**, *19*, 409.

(27) Liu, N.; Xu, Y.; Hossain, S.; Huang, N.; Coursolle, D.; Gralnick, J. A.; Boon, E. M. *Biochemistry* **2012**, *51*, 2087.

(28) Linares, J. F.; Gustafsson, I.; Baquero, F.; Martinez, J. L. Proc. Natl. Acad. Sci. U.S.A. 2006, 103, 19484.

(29) Hoffman, L. R.; D'Argenio, D. A.; MacCoss, M. J.; Zhang, Z. Y.; Jones, R. A.; Miller, S. I. *Nature* **2005**, *436*, 1171.

(30) Boehm, A.; Steiner, S.; Zaehringer, F.; Casanova, A.; Hamburger, F.; Ritz, D.; Keck, W.; Ackermann, M.; Schirmer, T.; Jenal, U. *Mol. Microbiol.* **2009**, *72*, 1500.

(31) Ahmed, N. A.; Petersen, F. C.; Scheie, A. A. Antimicrob. Agents Chemother. 2009, 53, 4258.

(32) He, M.; Ouyang, Z. M.; Troxell, B.; Xu, H. J.; Moh, A.; Piesman, J.; Norgard, M. V.; Gomelsky, M.; Yang, X. F. *PLoS Pathog.* **2011**, *7*, No. e1002133.

(33) Levi, A.; Folcher, M.; Jenal, U.; Shuman, H. A. *mBio* 2011, 2, No. e00316-10.

(34) Yang et al. Unpublished results.