In vivo Analysis of Cyclic di-GMP Cyclase and Phosphodiesterase Activity in Escherichia coli Using a Vc2 Riboswitch-based Assay

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#Contributed equally to this work

[Abstract] Cyclic di-guanosine monophosphate (c-di-GMP) is a ubiquitous second messenger that regulates distinct aspects of bacterial physiology. It is synthesized by diguanylate cyclases (DGCs) and hydrolyzed by phosphodiesterases (PDEs). To date, the activities of DGC and PDE are commonly assessed by phenotypic assays, mass spectrometry analysis of intracellular c-di-GMP concentration, or riboswitch-based fluorescent biosensors. However, some of these methods require cutting-edge equipment, which might not be available in every laboratory. Here, we report a new simple, convenient and cost-effective system to assess the function of DGCs and PDEs in E. coli. This system utilizes the high specificity of a riboswitch to c-di-GMP and its ability to regulate the expression of a downstream β-galactosidase reporter gene in response to c-di-GMP concentrations. In this protocol, we delineate the construction of this system and its use to assess the activity of DGC and PDE enzymes.

Keywords: Cyclic di-guanylate monophosphate (c-di-GMP), Diguanylate cyclase (DGC), Phosphodiesterase (PDE), Riboswitch, β-Galactosidase

[Background] Cyclic-di-GMP is an important and ubiquitous second messenger in bacteria, which regulates a variety of processes, such as motility-to-sessility transition, biofilm formation, virulence, and cell cycle progression (Römling et al., 2013). The GG(D/E)EF domain has diguanylate cyclase (DGC) activity and it is responsible for the synthesis of c-di-GMP from two GTPs, which is a two-step reaction with 5’-pppGpG as intermediate and two molecules of pyrophosphate as byproducts (Ryjenkov et al., 2005). Phosphodiesterases (PDE) with an EAL or an HD-GYP domain hydrolyze c-di-GMP into linear 5’-pGpG (Schmidt et al., 2005) and GMP (Ryan et al., 2006), respectively.

Several tools have been developed to monitor intracellular cyclic di-nucleotide levels and to identify proteins involved in cyclic di-nucleotide signaling, for example, protein-based fluorescence resonance energy transfer (FRET) biosensor (Christen et al., 2010), riboswitch-based fluorescent biosensor (Kellenberger et al., 2015), and riboswitch-based dual-fluorescence reporter (Zhou et al., 2016). However, these tools monitor altered fluorescence of reporters and require the access to flow cytometry or fluorescence microscopy. Here, we report the development of an alternative assay to monitor the intracellular c-di-GMP concentration, namely by monitoring the alteration in β-galactosidase
activity in agar-growing cells. For that, the Vc2 riboswitch (Sudarsan et al., 2008) is fused translationally to lacZY and integrated into the chromosome of E. coli strain TOP10. Vc2 is an ‘off’ riboswitch from Vibrio cholerae and thus down-regulates the expression of β-galactosidase when c-di-GMP is bound (Figure 1). The stable integration into the Tn7 attachment site in the chromosome of E. coli avoids copy number effects and eliminates the need to use an antibiotic resistance marker. Changes in c-di-GMP levels are subsequently translated to the alteration in β-galactosidase expression, which is reflected by the color change of the colony growing on an agar plate containing 5-Bromo-4-chloro-3-indolyl-β-D-galactopyranoside (X-gal). This assay can be used, for instance, to reveal the function of proteins under physiological condition and to assess the enzymatic activity of proteins that are challenging to be purified and tested in vitro. However, the Vc2-based assay described here is a qualitative assessment of the change in intracellular c-di-GMP concentration. Quantification is not crucial for a screening assay, but would be advantageous in, for example, measuring the activity of enzymes. We have demonstrated that the Vc2-based assay can be exploited to verify the activity of both DGCs and PDEs in vivo (El Mouali et al., 2017).

Figure 1. The principle of a riboswitch-based screening system. The Vc2 riboswitch is located upstream of the β-galactosidase open reading frame to control its expression in response to the variation in c-di-GMP concentration. When c-di-GMP is present in high abundance due to the overexpression of DGCs, the expression of β-galactosidase is down-regulated resulting in a white colony on an X-gal containing plate. In contrast, when PDEs are overexpressed, generating low intracellular c-di-GMP concentration, the colony is blue. In E. coli TOP10 wild type cells, there are residual amounts of c-di-GMP, resulting in a light blue colony (adapted from El Mouali et al., 2017).
Materials and Reagents

1. PCR tubes
2. Petri dish 92 x 16 mm (SARSTEDT, catalog number: 82.1472.001)
3. Syringe filters, 0.2 μm pore size (SARSTEDT, catalog number: 83.1826.001)
4. Microcentrifuge tubes of 1.5 ml (SARSTEDT, catalog number: 72.690.001)
5. Pipette tips (1,000 μl: SARSTEDT, catalog number: 70.762.100; 1-200 μl: Corning, catalog number: 4804; 0.1-10 μl: Gilson, catalog number: F171103)
6. Spectrophotometer cuvettes (SARSTEDT, catalog number: 67.742)
7. 3-part disposable HSW SOFT-JECT® syringes (5 ml: Henke Sass Wolf, catalog number: 5050.X00V0; 10 ml: Henke Sass Wolf, catalog number: 5100.X00V0)
8. Aluminum foil
9. Plasmids used in this study:

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>pGRG25</td>
<td>mTn7:MCS; Amp’</td>
<td>McKenzie and Craig, 2006</td>
</tr>
<tr>
<td>pBAD28</td>
<td>L-arabinose controlled expression vector; P_{BAD} promoter; AraC activator; pACYC184 ori; Amp’; Cm’</td>
<td>Guzman et al., 1995</td>
</tr>
<tr>
<td>pBAD30</td>
<td>L-arabinose controlled expression vector; P_{BAD} promoter; AraC activator; pACYC184 ori; Amp’</td>
<td>Guzman et al., 1995</td>
</tr>
<tr>
<td>pWJB30</td>
<td>pBAD30 with Xbal/HindIII-ligated adrA-6xHis fragment</td>
<td>Simm et al., 2004</td>
</tr>
<tr>
<td>pBADydeH</td>
<td>pBAD28 with HindIII/Xbal-ligated ydeH fragment</td>
<td>Jonas et al., 2008</td>
</tr>
<tr>
<td>pBADydeH-mut</td>
<td>mutagenized YdeH_{G206A/G207A}</td>
<td>Jonas et al., 2008</td>
</tr>
<tr>
<td>p4551</td>
<td>pBAD30 with SacI/HindIII-ligated STM4551-6xHis fragment</td>
<td>Ahmad et al., 2011</td>
</tr>
<tr>
<td>p4551_{E267A}</td>
<td>mutagenized STM4551_{E267A}</td>
<td>Ahmad et al., 2011</td>
</tr>
<tr>
<td>pSRKGM</td>
<td>expression vector containing the P_{lac} promoter; LacI\textsuperscript{q} repressor, lacZα\textsuperscript{+}:MCS, pBBR1 ori; Gm’</td>
<td>Khan et al., 2008</td>
</tr>
<tr>
<td>pSTM3611</td>
<td>pSRKGM with Ndel/Nhel-ligated STM4551-6xHis fragment</td>
<td>El Mouali et al., 2017</td>
</tr>
<tr>
<td>pSTM3611_{E136A}</td>
<td>mutagenized STM3611_{E136A}</td>
<td>El Mouali et al., 2017</td>
</tr>
<tr>
<td>Vc2-pRS414</td>
<td>-348 nt to +21 nt of COG3070 ORF cloned in pRS414 vector as an EcoRI-BamHI fragment</td>
<td>Sudarsan et al., 2008</td>
</tr>
<tr>
<td>pGRG25:Vc2lacZY</td>
<td>pGRG25 with Pael/NotI-ligated Vc2lacZY fragment</td>
<td>El Mouali et al., 2017</td>
</tr>
</tbody>
</table>
10. Calcium chloride competent *E. coli* TOP10 cells (Hanahan, 1983)

11. Primers used in this study (ordered from Sigma-Aldrich):

<table>
<thead>
<tr>
<th>Name</th>
<th>Sequence</th>
</tr>
</thead>
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<tr>
<td>Vc2lacZY_F _PacI</td>
<td>GCGTTAATTAATAACGCTATATTGAAAGCTTG*</td>
</tr>
<tr>
<td>Vc2lacZY_R_NotI</td>
<td>GATCAAGCAGCCTTAAGCGACTTCATTCACCTG</td>
</tr>
<tr>
<td>pGRGcontrol_F</td>
<td>CACTTATCTGGTGTGTCGACACT</td>
</tr>
<tr>
<td>Vc2lacZY_R</td>
<td>CTGCAAGCAGATTAAGTTGG</td>
</tr>
<tr>
<td>pGRGcontrol_F</td>
<td>GATGCTGTGGCGCGAAGCTGT</td>
</tr>
<tr>
<td>pGRGcontrol_R</td>
<td>GATGACGGTTTGTCCATGGG</td>
</tr>
</tbody>
</table>

*a The restriction sites are underlined.*

12. GenElute Plasmid Miniprep Kit (Sigma-Aldrich, catalog number: PLN350)

13. Sterile distilled, deionized water (diH2O)


15. dNTP mix (10 mM each) (Thermo Fisher Scientific, Thermo Scientific™, catalog number: R0192)

16. Agarose (Sigma-Aldrich, catalog number: A9539)

17. SmartLadder for DNA: 200-10,000 bp (Eurogentec, catalog number: MW-1700-10)

18. GelRed™ nucleic acid gel stain (Biotium, catalog number: 41003)

19. *NotI* restriction enzyme (New England Biolabs, catalog number: R0189S)

20. *PacI* restriction enzyme (New England Biolabs, catalog number: R0547S)

21. PCR purification kit (QIAGEN, catalog number: 28106)

22. Rapid DNA ligase kit (Roche Diagnostics, catalog number: 10716359001)

23. Taq DNA polymerase with standard Taq buffer (New England Biolabs, catalog number: M0273S)

24. Sodium chloride (Sigma-Aldrich, catalog number: S7653)

25. Tryptone (BD, Bacto™, catalog number: 211705)

26. Yeast extract (BD, Bacto™, catalog number: 212750)

27. Agar (BD, Difco™, catalog number: 281230)

28. Tris base (Sigma-Aldrich, catalog number: 93350)

29. Glacial acetic acid (Sigma-Aldrich, catalog number: ARK2183)

30. EDTA (Sigma-Aldrich, catalog number: E9884)

31. Ampicillin sodium salt (Sigma-Aldrich, catalog number: A9518)

32. Gentamicin sulfate salt (Sigma-Aldrich, catalog number: G3632)

33. L-(-)-arabinose (Sigma-Aldrich, catalog number: A3256)

34. Isopropyl β-D-1-thiogalactopyranoside (IPTG) (Sigma-Aldrich, catalog number: I6758)

35. Dimethyl sulfoxide (Sigma-Aldrich, catalog number: D8418)

36. 5-Bromo-4-chloro-3-indolyl-β-D-galactopyranoside (X-gal) (Roche Diagnostic, catalog number: 10703729001)

37. LB (Luria-Bertani) medium (see Recipes)

38. LB (Luria-Bertani) agar (see Recipes)
39. TAE buffer (see Recipes)
40. 100 mg/ml ampicillin stock (see Recipes)
41. 30 mg/ml gentamicin stock (see Recipes)
42. 100 mM IPTG stock (see Recipes)
43. 20 mg/ml X-gal stock (see Recipes)

**Equipment**

1. Erlenmeyer flasks (50 ml/100 ml/250 ml)
2. Pipettes [e.g., PIPETMAN® P20 (Gilson, catalog number: F123600), P200 (Gilson, catalog number: F123601), or P1000 (Gilson, catalog number: F123602)]
3. SureCycler 8800 thermal cycler (Agilent Technologies, model: SureCycler 8800, catalog number: G8800A)
5. Mini-Sub® cell GT Horizontal Electrophoresis System (Bio-Rad Laboratories, catalog number: 1704406)
6. 15-well comb (Bio-Rad Laboratories, catalog number: 1704464)
7. Sub-Cell GT UV-Transparent Mini-Gel Tray (Bio-Rad Laboratories, catalog number: 1704436)
8. Electrophoresis power supply
9. Heraeus® microbiological incubator (Thermo Fisherer Scientific, Thermo Scientific™, model: Heraeus® microbiological incubator)
10. Multitron Standard shaker (INFORS HT)
11. Gel Doc™ XR+ Gel Documentation System (Bio-Rad Laboratories, catalog number: 1708195)
12. BioPhotometer (Eppendorf, model: 6131)
14. Thermomixer compact (Eppendorf, model: 5350)
15. Digital camera

**Procedure**

Practically, the workflow can be divided into two parts, as shown in Figure 2. The first part is to construct the reporter strain, which takes approximately 12-15 days (Steps 1-3 of the protocol). The second part is to assess the activity of DGC and PDE by the reporter strain (Steps 4-5). This will take 7-9 days.
**Figure 2. The overview of the workflow.** The procedure is divided into two parts: 1) construction of the reporter strain and 2) assessment of c-di-GMP cyclase or phosphodiesterase activity by the reporter. It takes approximately 12-15 days to construct the reporter strain, and another 7-9 days to assess the target.

1. Amplify Vc2 and *lacZ*Y by PCR:
   a. The template for the PCR is Vc2-pRS414 (kindly received from Prof. R. R. Breaker, Yale University), in which the fragment containing riboswitch along with its native promoter was cloned in frame to the *lacZ* reporter gene (Sudarsan et al., 2008).
   b. The primers used are:
      - Vc2lacZ\_Y\_\_F\_Paci: GCGTTAATTAAATAACGCCTATATTGAAAGCTTG
      - Vc2lacZ\_Y\_\_R\_NotI: GATCAGCGGCCGCTTAAGCGACTTCATTCACCTG
      The restriction sites PacI and NotI are underlined.
   c. Prepare PCR reaction mix:
      
      | Component | Volume | Final concentration |
      |-----------|--------|---------------------|
      | 5x Phusion HF buffer | 10 μl | 1x |
      | 10 mM dNTP | 1 μl | 200 μM each |
      | 10 μM forward primer | 2.5 μl | 0.5 μM |
      | 10 μM reverse primer | 2.5 μl | 0.5 μM |
      | Phusion DNA polymerase | 0.5 μl | 0.02 U/μl |
      | Vc2-pRS414 plasmid | X μl | 1 pg-10 ng per 50 μl reaction volume |
      | H₂O | Add to 50 μl |

d. Thermocycling conditions:
   - 98 °C for 30 sec \( \times \) 1 cycle
   - 98 °C for 10 sec
   - 62 °C for 30 sec \( \times \) 35 cycles
   - 72 °C for 150 sec
   - 72 °C for 10 min \( \times \) 1 cycle
   - Hold at 4 °C
   e. Check the PCR products by gel electrophoresis using 1% TAE agarose gel. The expected size is 4,733 bp.
2. Clone Vc2lacZ\_Y\_pGRG25 vector:
a. Double digest the PCR product Vc2lacZY and the plasmid pGRG25 with PacI and NotI at 37 °C overnight. To reduce star activity, it is advisable to use the fewest units possible to achieve the digestion. After digestion, it is necessary to check the size of products using agarose gel electrophoresis. The enzymes are inactivated at 65 °C for 20 min.

b. Remove the restriction enzymes using PCR purification kit. The concentration of digested Vc2lacZY and pGRG25 are measured by NanoDrop.

c. Ligate pGRG25 and Vc2lacZY of molar ratio 1:1 at 12 °C overnight using T4 DNA ligase following the manufacturer’s instruction. To verify that the vector is completely cut, ligate the digested vector alone. If the vector is not cut by the two restriction enzymes efficiently, re-circulation will happen and colonies will appear after transformation.

d. Transform the ligation products into chemically competent TOP10 cells and select for the recombinant plasmids on LB agar plate containing 100 μg/ml ampicillin. The pGRG25 vector is temperature-sensitive, so the plate should be incubated at 32 °C or even lower temperature.

e. Verify the recovered colonies by colony PCR. First, prepare the master mix for Taq DNA polymerase with standard Taq buffer (from New England Biolabs) following manufacturer's instruction. Then 25 µl aliquots are added to the PCR tubes. Sterile pipette tips are used to pick single colonies and mix with reaction buffer. An initial 5 min denaturation at 95 °C is recommended for colony PCR with Taq DNA polymerase. The expected size is 518 bp. The primers used are:

   pGRG25control_F: CACTTATCTGGTTGGTCGACACT
   Vc2lacZY_R: CTGCAAGGCGATTAAGTTGG

   The details for the colony PCR can be found in the protocol by Azevedo et al. (2017).

f. Confirm the sequence integrity of the riboswitch in the positive clones by DNA sequencing using the same primers for colony PCR.

3. Integrate Vc2lacZY into the Tn7 attachment site of E. coli chromosome (Note 1):

a. Streak positive colonies on LB agar plate supplemented with 100 μg/ml ampicillin and incubate the cells at 32 °C.

b. Inoculate approximately 10 colonies into 3 ml of ampicillin-free LB medium at 32 °C, 200 rpm. Add 0.1% L-arabinose (w/v) to facilitate the incorporation of Vc2lacZY into E. coli Tn7 attachment site. The integration results in a new strain E. coli TOP10 attTn7::Vc2lacZY.

The details for the colony PCR can be found in the protocol by Azevedo et al. (2017).

c. Dilute the overnight culture by a factor of 10^8 and plate 50 µl of the diluted culture on LB plate to acquire single colonies. Incubate the plate at 42 °C to block the replication of the temperature-sensitive pGRG25 vector.

d. Streak individual colonies on LB plate and repeat the incubation at 42 °C to make sure that the plasmid is lost.

e. Confirm the integration of Vc2lacZY into the chromosome by colony PCR using Taq DNA polymerase. The primers used are:

   pGRGcontrol_F: GATGCTGGTGCGAAGCTGT
4. Assess the activity of diguanylate cyclases by Vc2-based screening: take the established DGC AdrA, YdeH and STM4551 as examples
   a. Prepare calcium chloride competent *E. coli* TOP10 attTn7::Vc2lacZY cells.
   b. Transform pBAD vectors (Note 2) harboring DGC (AdrA, YdeH and STM4551) and catalytically inactive mutants (STM4551E267A and YdeH-mut) as well as empty vectors into the chemically competent cells by heat shock at 42 °C for 40 sec. Add 1 ml LB medium free of antibiotics into the microcentrifuge tubes and incubate the tubes at 37 °C, 200 rpm for 1 h to let the cells recover.
   c. Plate 200 μl cells on LB agar plates containing 100 μg/ml ampicillin and incubate the plates at 37 °C overnight.
   d. Streak single colonies on ampicillin-containing LB agar plates.
   e. Pick one colony for each strain and culture the colony in 5 ml of LB medium with 100 μg/ml ampicillin at 37 °C, 200 rpm overnight.
   f. Measure the OD600 of overnight culture and adjust the OD600 to 0.1.
   g. Grow cells at 37 °C, 200 rpm. When OD600 reaches 0.6-1.0, spot 3 μl culture on LB plates with the addition of 100 μg/ml ampicillin, 80 μg/ml X-gal, 0.1% (w/v) L-arabinose (Note 3). It is important that cells with DGC, corresponding mutant, and vector control have similar OD600.
   h. Incubate the plates at 28 °C and monitor the color of colonies up to 72 h. A DGC activity will lead to a white colony indicative of high c-di-GMP levels. Take pictures of the plate using a digital camera.

5. Investigate the activity of PDE: take the established PDE STM3611 as an example
   a. The intracellular concentration of c-di-GMP is low in TOP10 cells, therefore Vc2lacZY-harboring cells have a basal level of β-galactosidase activity resulting in a blue colony, which makes it impossible to distinguish the color further enhanced by phosphodiesterase. The activity of PDE can, however, be reliably investigated by co-expression of a DGC. Therefore, pWJB30 (DGC AdrA cloned into pBAD30) and pSTM3611 (PDE STM3611 cloned into pSRKGm) (Notes 2 and 4) are co-transformed into chemically competent TOP10::Vc2lacZY cells. As controls, empty vectors are co-transformed or in combination with the DGC/PDE. The combinations are listed in Table 1.
Table 1. The combination of plasmids co-transformed into chemically competent TOP10::Vc2lacZY cells

<table>
<thead>
<tr>
<th>No.</th>
<th>Plasmid for PDE</th>
<th>Plasmid for DGC</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>pSRKGm (vector control)</td>
<td>pBAD30 (vector control)</td>
</tr>
<tr>
<td>2</td>
<td>pSRKGm</td>
<td>pWJB30 (AdrA in pBAD30)</td>
</tr>
<tr>
<td>3</td>
<td>pSTM3611 (STM3611 in pSRKGm)</td>
<td>pBAD30</td>
</tr>
<tr>
<td>4</td>
<td>pSTM3611</td>
<td>pWJB30</td>
</tr>
<tr>
<td>5</td>
<td>pSTM3611E136A (catalytic inactive STM3611 in pSRKGm)</td>
<td>pWJB30</td>
</tr>
</tbody>
</table>

b. Plate cells on LB agar plates containing both ampicillin (100 μg/ml) and gentamicin (30 μg/ml). The plates are incubated at 37 °C overnight.

c. Streak the positive colonies on antibiotics-containing LB agar plates.

d. Inoculate at least three individual colonies per strain in 5 ml of LB medium containing both ampicillin and gentamicin. Grow cells at 37 °C, 200 rpm overnight.

e. Measure OD600 of overnight culture and dilute the culture to OD600 of 0.1 with LB medium containing antibiotics. Let the cells grow to exponential phase (OD600 ~0.6). It is critical that all samples have similar OD600 values.

f. Spot 3 μl of the cultures on LB agar plates containing ampicillin (100 μg/ml), gentamicin (30 μg/ml), IPTG (0.75 mM), L-arabinose (0.003%, w/v) and X-Gal (80 μg/ml) (Note 3).

g. Incubate the plate at 28 °C and record the color of the colonies up to 72 h. While expression of the DGC alone leads to white colonies, co-expression of an active PDE will lead to blue colonies.

h. Take pictures of the plate using a digital camera.

Data analysis

1. Assessment of the activity of DGC by the Vc2-based assay: after spotted on X-gal containing plates, the color of colonies is monitored periodically. Figure 3 shows the color after 24 h incubation at 28 °C. It is explicit that cells expressing DGCs have whiter color due to the riboswitch inhibition of β-galactosidase expression upon high c-di-GMP levels compared with the corresponding mutant and vector control.

Figure 3. Detection of established diguanylate cyclases by the Vc2-based assay. DGC AdrA, STM4551 and catalytic inactive STM4551E267A were cloned into pBAD30 generating pWJB30, p4551 and p4551E267A, respectively. Another DGC YdeH and its mutant YdeHG206AG207A were cloned into pBAD28 resulting in pBAD28ydeH and pBAD28ydeH-mut. Cells expressing functional DGC (AdrA/STM4551/YdeH) are white, while cells with vector
control or catalytically inactive mutant (STM4551E267A/YdeH2206AG207A) are blue. Growth was at 28 °C for 24 h on LB agar plates supplemented with 100 μg/ml ampicillin, 80 μg/ml X-gal and 0.1% arabinose (w/v).

2. Assessment of the enzymatic activity of a PDE by the Vc2-based assay: Figure 4 shows the color of colonies after incubation at 28 °C for 48 h. Cells expressing the DGC AdrA alone (row 2) or co-expressing AdrA and catalytically inactive PDE STM3611E136A (row 5) are white. However, cells are blue when wild type STM3611 is co-expressed with AdrA (row 4), which illustrates the enzymatic activity of STM3611 and the decrease of the intracellular c-di-GMP concentration upon expression.

![Figure 4](image)

**Figure 4. Detection of phosphodiesterase activity of PDE STM3611 by the Vc2-based assay.** Phosphodiesterase STM3611 and catalytic inactive STM3611E136A were cloned into pSRKGm generating pSTM3611 and pSTM3611E136A, respectively. Diguanylate cyclase AdrA was cloned into pBAD30 resulting in pWJB30. Cells expressing the DGC AdrA alone have a white color. Upon co-expressing the PDE STM3611 with AdrA, the color turns to blue indicating low c-di-GMP level. However, cells expressing the DGC AdrA with catalytically inactive PDE STM3611E136A are white. Growth was at 28 °C for 48 h on LB agar plate supplemented with 100 μg/ml ampicillin, 30 μg/ml gentamicin, 80 μg/ml X-gal, 0.003% arabinose (wt/vol), and 0.75 mM IPTG (adapted from El Mouali et al., 2017).

**Notes**

1. The protocol for the integration of target genes into the *E. coli* chromosomal Tn7 attachment site is delineated by G. J. McKenzie and N. L. Craig (McKenzie and Craig, 2006).

2. We cloned diguanylate cyclases into pBAD28/pBAD30 vectors and phosphodiesterases into the pSRKGm vector. Other expression vectors can also be used depending on the availability.
3. Different L- (+)-arabinose and IPTG concentrations need to be tested to achieve the most discriminating color contrast between diguanylate cyclase, phosphodiesterase and control.

4. It is important that diguanylate cyclase and phosphodiesterase are cloned into compatible vectors containing different antibiotic resistance markers and compatible origins of replication.

5. The specificity of the Vc2-based riboswitch towards other cyclic di-nucleotides has not been tested in this work.

**Recipes**

1. LB (Luria-Bertani) medium for 400 ml
   - 4 g NaCl
   - 4 g tryptone
   - 2 g yeast extract
   - Add dH2O to 400 ml
   - Autoclave at 121 °C for 15 min

2. LB (Luria-Bertani) agar for 400 ml
   - 4 g NaCl
   - 4 g tryptone
   - 2 g yeast extract
   - 6 g agar
   - Add dH2O to 400 ml
   - Autoclave at 121 °C for 15 min
   - Pour into Petri dishes (add antibiotics, X-gal, L-arabinose, or IPTG if required)
   - Let cool completely and store at 4 °C

3. TAE buffer (50x stock solution)
   - Dissolve 242 g Tris base in water
   - Add 57.1 ml glacial acetic acid
   - Add 100 ml of 0.5 M EDTA (pH 8.0)
   - Add water up to 1 L

4. 100 mg/ml ampicillin stock
   - Dissolve 500 mg ampicillin sodium salt in 5 ml deionized water
   - Filter the solution with 0.2 μm sterile syringe filter
   - Make 1 ml aliquot of solution into sterile microcentrifuge tubes
   - Store at -20 °C

5. 30 mg/ml gentamicin stock
   - Dissolve 150 mg gentamicin sulfate salt in 5 ml deionized water
   - Filter the solution with 0.2 μm sterile syringe filter
   - Make 1 ml aliquot of solution into sterile microcentrifuge tubes
   - Store at -20 °C
6. 100 mM IPTG stock
   - Dissolve 1.191 g IPTG in 5 ml deionized water
   - Filter the solution with 0.2 µm sterile syringe filter
   - Make 1 ml aliquot of solution into sterile microcentrifuge tubes
   - Store at -20 °C

7. 20 mg/ml X-gal stock
   - Dissolve 100 mg X-gal in 5 ml DMSO
   - Wrap the tube with aluminum foil to protect from light
   - Store at -20 °C
   - The solution is stable for 6-12 months at -20 °C

Acknowledgments

The authors would like to thank Prof. Dr. Ronald R. Breaker for sharing Vc2-pRS414 riboswitch construct. Prof. Ute Römling conceived this study. Ying Liu developed the assay and assessed DGC activity. Hyunhee Kim developed the assay to assess PDE activity. This work was supported by the Swedish Research Council for Natural Sciences and Engineering (grant 621-2013-4809) and the Karolinska Institutet. This modified protocol is based on a previously published work (El Mouali et al., 2017). The authors declare no conflicts of interest or competing interests.

References


