

**Phosphodiesterase Activity of *Ec* DOS, a Heme-regulated Enzyme
from *Escherichia coli*, toward 3',5'-Cyclic Diguanylic Acid Is Obviously
Enhanced by O₂ and CO Binding**

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3',5'-Cyclic diguanylic acid (c-di-GMP) is emerging as an important bacterial intracellular signaling molecule. Both the Fe(III) and Fe(II) heme complexes of *Ec* DOS, a heme-bound PAS sensor enzyme from *Escherichia coli*, have phosphodiesterase activity toward c-di-GMP (27 and 61 min⁻¹, respectively) and that the activity of the Fe(II) heme complex is obviously enhanced by the binding of either O₂ (126 min⁻¹) or CO (143 min⁻¹). Therefore, *Ec* DOS appears to be a novel type of gas sensor enzyme.

The heme-bound domain of *Ec* DOS, a heme-sensor enzyme from *Escherichia coli*, was first cloned by the Gilles-Gonzalez group.¹ Based on its physicochemical characteristics and comparison with the direct oxygen sensor enzyme, FixL, which has a heme-bound PAS sensor domain, they predicted that *Ec* DOS acts as a direct oxygen sensor enzyme and that O₂ binding to the heme iron of this enzyme should suppress its phosphodiesterase activity. In separate experiments, this group demonstrated that the full-length enzyme is a cAMP phosphodiesterase.^{2,3} This and other groups have also characterized the physicochemical and structural properties of the isolated heme-bound PAS domain.⁴⁻⁹ They found that the cAMP phosphodiesterase activity is regulated by the heme redox state: it is active when the heme is in the Fe(II) state and inactive with heme in the Fe(III) state.^{2,3} Also, binding of CO or NO to the Fe(II) complex makes the

enzyme inactive toward cAMP.² Profound structural changes in the heme-bound PAS accompanying the heme redox change may explain the heme redox-dependent activity toward cAMP.⁶ Knockout of *Ec* DOS in *E. coli* causes the level of cAMP to increase 27-fold compared with the wild type and induces alterations in cell morphology, suggesting that cAMP is a substrate of *Ec* DOS in vivo and that *Ec* DOS is involved in cell differentiation.^{10,11}

Recently, 3',5'-cyclic diguanylic acid (c-di-GMP) has emerged as an important bacterial intracellular signaling molecule associated with motility, virulence, intercellular interaction, and cellulose production.¹¹⁻¹⁸ The C-terminal domain of *Ec* DOS has a GGDEF domain, which is predicted to have diguanylate cyclase activity (conversion of GTP to c-di-GMP), and an EAL domain, which should act as a c-di-GMP-specific phosphodiesterase (conversion of c-di-GMP to l-di-GMP). Although, the isolated C-terminal domain has c-di-GMP-specific phosphodiesterase activity,¹⁵ the effect of the heme-bound N-terminal domain of full-length *Ec* DOS has not been reported.

We examined the catalytic activity of *Ec* DOS toward c-di-GMP by HPLC, using NADPH as a standard. In the absence of *Ec* DOS, there was no difference in the HPLC patterns under aerobic and anaerobic conditions (Figure 1a). Upon the addition of Fe(III) *Ec* DOS, there were marked changes in the HPLC pattern; specifically, there was a rapid decrease in the c-di-GMP peak and a concomitant increase in the l-di-GMP peak. This

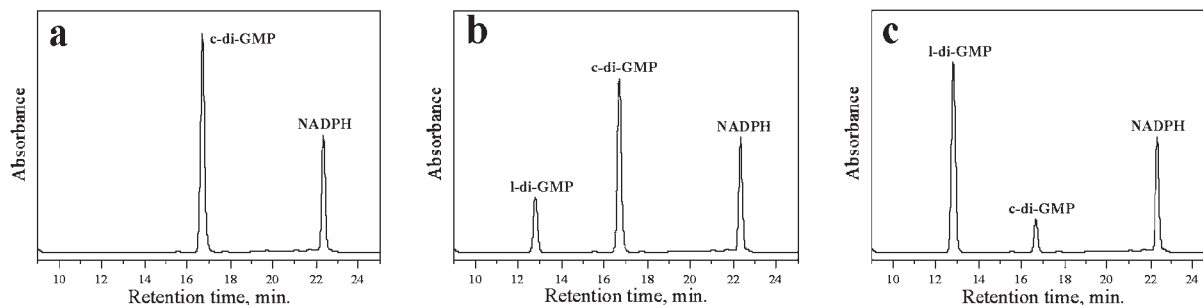


Figure 1. HPLC patterns of (a) c-di-GMP (100 μ M) and NADPH (standard) in the absence of *Ec* DOS; (b) the reaction mixture containing l-di-GMP, c-di-GMP, and NADPH 3 min after addition of *Ec* DOS; and (c) the CO-saturated reaction solution containing l-di-GMP, c-di-GMP, and NADPH 3 min after addition of *Ec* DOS. To avoid any differences in experimental conditions, all experiments were conducted in a glove box at an oxygen concentration of less than 10 ppm. The Fe(III) and Fe(II) enzymes were obtained by mixing *Ec* DOS with ferricyanide and sodium dithionite, respectively, followed by removal of the excess redox reagents by Sephadex G-25 column chromatography. Phosphodiesterase activity was assayed at 37 °C for 3 min in a reaction mixture containing 50 mM Tris-HCl (pH 8.5), 50 mM NaCl, 5 mM MgCl₂, 0.1 mM c-di-GMP, and 0.2 μ M *Ec* DOS. The reaction was stopped by addition of CaCl₂ (final concentration, 10 mM) and adjusted to 0.1 mM β -NADPH (standard). The mixture was then filtered through a 0.45- μ m filter, and the reaction samples (10 μ L) were injected into a LUNA 5 μ C18 (2) column (15 \times 4.6 cm; Phenomenex, Torrance, CA, U.S.A.) and analyzed using a Prominence HPLC system (Shimadzu, Kyoto, Japan) with detection at 254 nm. The following buffers were used in the gradient program: buffer A (100 mM KH₂PO₄ with 4 mM tetrabutyl ammonium hydrogen sulfate [pH 6.0]) and buffer B (75% buffer A/25% methanol). The gradient was delivered at a flow rate of 0.7 mL min⁻¹ according to the following program: 0.0 min, 40% B/60% A; 15.0 min, 100% B; 20.0 min, 100% B; and 21.0 min, 40% B/60% A.

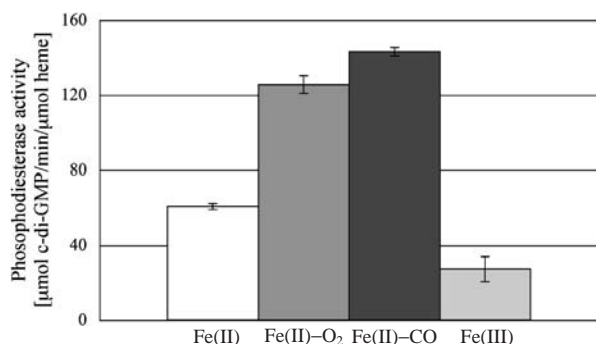


Figure 2. The rates of substrate hydrolysis under various conditions. From left to right, activities of Fe(II), Fe(II)-O₂, Fe(II)-CO, and Fe(III) forms of *Ec* DOS. The activity of the Fe(II) complex of *Ec* DOS was enhanced by CO and O₂ binding. Reaction rates represent means of six determinations.

HPLC pattern is the same as previously reported.^{14–16} Figure 1b shows the HPLC pattern obtained after a 3-min incubation at 37 °C with Fe(III) *Ec* DOS. During this 3-min reaction, the turnover number for the Fe(III) enzyme was 27 min⁻¹. Addition of Fe(II) *Ec* DOS gave similar time-dependent HPLC pattern, with a turnover number of 61 min⁻¹. This two-fold increase in activity for the Fe(II) enzyme contrasts with the enzyme's cAMP phosphodiesterase activity, which is strictly heme-redox dependent and inactive in the Fe(III) state.^{2,3}

We next examined the effects of O₂ and CO binding on the Fe(II) complex. Surprisingly, we found that the catalytic activity was obviously higher for the O₂- and CO-bound *Ec* DOS enzymes (126 and 143 min⁻¹, respectively; Figures 1c and 2). Because the rate of autooxidation of the full-length wild-type enzyme is slow (0.015 min⁻¹), the optical absorption spectra for the O₂-bound Fe(II) enzyme was stable during the course of activity measurement. Only a very small GMP peak was observed even after a 45-min reaction, suggesting that the activity toward 1-di-GMP was very low under the present conditions.

Several O₂- and CO-response heme-sensor enzymes and proteins (FixL,^{19,20} AXPDEA1,²¹ cystathion β-synthase,²² heme-bound neuronal PAS protein 2 (NPAS2),²³ and CoxA^{24,25}) are known. The functions of most of these are suppressed by O₂ and CO binding, except for CoxA. Therefore, the present study shows that the Fe(II) complex of *Ec* DOS is the first heme-sensor enzyme with a heme-bound PAS domain whose activity is enhanced by the binding of O₂ and CO. In particular, AXPDEA1 has both GGDEF and EAL domains the same as *Ec* DOS and the sequence homology of the C-terminal catalytic domains between AXPDEA1 and *Ec* DOS is high, 30% identical and 50% homologous.^{1,21} However, both enzymes exhibited the opposite ligand effects in terms of the PDE activity toward c-di-GMP.

Further studies using physical methods are needed to elucidate the unique intramolecular signaling activated in *Ec* DOS by O₂ and CO binding.²⁶

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References

- 1 V. M. Delgado-Nixon, G. Gonzalez, M. A. Gilles-Gonzalez, *Biochemistry* **2000**, *39*, 2685.
- 2 Y. Sasakura, S. Hirata, S. Sugiyama, S. Suzuki, S. Taguchi, M. Watanabe, T. Matsui, I. Sagami, T. Shimizu, *J. Biol. Chem.* **2002**, *277*, 23821.
- 3 T. Yoshimura, I. Sagami, Y. Sasakura, T. Shimizu, *J. Biol. Chem.* **2003**, *278*, 53105.
- 4 A. Sato, Y. Sasakura, S. Sugiyama, I. Sagami, T. Shimizu, Y. Mizutani, T. Kitagawa, *J. Biol. Chem.* **2002**, *277*, 32650.
- 5 S. Taguchi, T. Matsui, J. Igarashi, Y. Sasakura, Y. Araki, O. Ito, S. Sugiyama, I. Sagami, T. Shimizu, *J. Biol. Chem.* **2004**, *279*, 3340.
- 6 H. Kurokawa, D. S. Lee, M. Watanabe, I. Sagami, B. Mikami, C. S. Raman, T. Shimizu, *J. Biol. Chem.* **2004**, *279*, 20186.
- 7 H. J. Park, C. Suquet, J. D. Satterlee, C. H. Kang, *Biochemistry* **2004**, *43*, 2738.
- 8 Y. Sasakura, T. Yoshimura-Suzuki, H. Kurokawa, T. Shimizu, *Acc. Chem. Res.* **2006**, *39*, 37.
- 9 G. Gonzalez, E. Dioum, C. M. Bertolucci, T. Tomita, M. Ikeda-Saito, M. R. Cheesman, N. J. Watmough, M. A. Gilles-Gonzalez, *Biochemistry* **2002**, *41*, 8414.
- 10 T. Yoshimura-Suzuki, I. Sagami, N. Yokota, H. Kurokawa, T. Shimizu, *J. Bacteriol.* **2005**, *187*, 6678.
- 11 M. M. Méndez-Ortiz, M. Hyodo, Y. Hayakawa, J. Membrillo-Hernández, *J. Biol. Chem.* **2006**, *281*, 8090.
- 12 R. Simm, M. Morr, A. Kader, M. Nimtz, U. Römling, *Mol. Microbiol.* **2004**, *53*, 1123.
- 13 L. R. Hoffman, D. A. D'Argenio, M. J. MacCoss, Z. Zhang, R. A. Jones, S. I. Miller, *Nature* **2005**, *436*, 1171.
- 14 D. A. Ryjenkov, M. Tarutina, O. V. Moskvina, M. Gomelsky, *J. Bacteriol.* **2005**, *187*, 1792.
- 15 A. J. Schmidt, D. A. Ryjenkov, M. Gomelsky, *J. Bacteriol.* **2005**, *187*, 4774.
- 16 M. Christen, B. Christen, M. Folcher, A. Schauerte, U. Jena, *J. Biol. Chem.* **2005**, *280*, 30829.
- 17 Tamayo, A. D. Tischler, A. Camilli, *J. Biol. Chem.* **2005**, *280*, 33324.
- 18 H. Kulesekara, V. Lee, A. Brenic, N. Liberati, J. Urbach, S. Miyata, D. G. Lee, A. N. Neely, M. Hyodo, Y. Hayakawa, F. M. Ausubel, S. Lory, *Proc. Natl. Acad. Sci. U.S.A.* **2006**, *103*, 2839.
- 19 E. H. S. Souana, G. Gonzalez, M. A. Gilles-Gonzalez, *Biochemistry* **2005**, *44*, 15359.
- 20 A. Tanaka, H. Nakamura, Y. Shiro, H. Fujii, *Biochemistry* **2006**, *45*, 2515.
- 21 A. L. Chang, J. R. Tuckman, G. Gonzalez, R. Mayer, H. Weinhouse, G. Volman, D. Amikam, M. Benziman, M. A. Gilles-Gonzalez, *Biochemistry* **2001**, *40*, 3420.
- 22 R. Banerjee, C. Zou, *Arch. Biochem. Biophys.* **2005**, *433*, 144.
- 23 E. M. Dioum, J. Rutter, J. R. Tuckerman, G. Gonzalez, M. A. Gilles-Gonzalez, S. L. McKnight, *Science* **2002**, *298*, 2385.
- 24 H. Youn, R. L. Kerby, G. P. Roberts, *J. Biol. Chem.* **2004**, *279*, 45744.
- 25 S. Aono, *Acc. Chem. Res.* **2003**, *36*, 825.
- 26 T. Uchida, T. Kitagawa, *Acc. Chem. Res.* **2005**, *38*, 662.