

Extracellular Ca2+ transients affect poly-(R)-3-hydroxybutyrate regulation by the AtoS-AtoC system in E. coli

Marina C Theodorou, Ekaterini Tiligada, Dimitrios A Kyriakidis

▶ To cite this version:

Marina C Theodorou, Ekaterini Tiligada, Dimitrios A Kyriakidis. Extracellular Ca2+ transients affect poly-(R)-3-hydroxybutyrate regulation by the AtoS-AtoC system in E. coli. Biochemical Journal, 2009, 417 (3), pp.667-672. 10.1042/BJ20081169. hal-00479043

HAL Id: hal-00479043 https://hal.science/hal-00479043

Submitted on 30 Apr 2010

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

Extracellular Ca²⁺ transients affect poly-(R)-3-hydroxybutyrate regulation by the AtoS-AtoC system in *E. coli*

Marina C. THEODOROU*, Ekaterini TILIGADA† and Dimitrios A. KYRIAKIDIS*'[‡],¹

* Laboratory of Biochemistry, Department of Chemistry, Aristotle University of Thessaloniki, GR-54124 Thessaloniki, Greece

[†] Department of Pharmacology, Medical School, University of Athens, M. Asias 75, GR-11527 Athens, Greece

‡ National Hellenic Research Foundation, Vas. Constantinou 48, GR-11635 Athens, Greece

Short title: Ca²⁺ affects cPHB regulation by the AtoS-AtoC system

¹Correspondence to: Prof. D.A. Kyriakidis Laboratory of Biochemistry Department of Chemistry Aristotle University of Thessaloniki GR-54124 Thessaloniki Tel: +302310997771 Fax: +302310997689 email: kyr@chem.auth.gr

Abstract

Escherichia coli is exposed to wide extracellular concentrations of Ca^{2+} , while the cytosolic levels of the ion are subject to stringent control and are implicated in many physiological functions. This study shows that extracellular Ca^{2+} controls complexed poly-(R)-3-hydroxybutyrate (cPHB) biosynthesis through the AtoS-AtoC two component system. Maximal cPHB accumulation was observed at higher $[Ca^{2+}]e$ in AtoS-AtoC expressing *E. coli* compared to their $\Delta atoSC$ counterparts, in both cytosolic and membrane fractions. The reversal of EGTA-mediated down-regulation of cPHB biosynthesis by the addition of Ca^{2+} and Mg²⁺ was under the control of the AtoS-AtoC system. Moreover, the Ca^{2+} channel blocker verapamil reduced total and membrane-bound cPHB levels, the inhibitory effect being circumvented by Ca^{2+} addition only in *atoSC*⁺ bacteria. Histamine and compound 48/80 affected cPHB accumulation in a $[Ca^{2+}]e$ -dependent manner directed by the AtoS-AtoC system. In conclusion, these data provide evidence for the involvement of external Ca^{2+} on cPHB synthesis regulated by the AtoS-AtoC two-component system, thus linking Ca^{2+} with a signal transduction system, most probably through a transporter.

Key words

AtoS-AtoC two-component system, *atoDAEB*, calcium, calcium channel blockers, histamine, poly-(R)-3-hydroxybutyrate

Abbreviations used: $[Ca^{2+}]e$, extracellular Ca^{2+} concentration; cPHB, complexed poly-(R)-3hydroxybutyrate; C48/80, compound 48/80; TCS, two-component system; URR, upstream regulatory region

THIS IS NOT THE VERSION OF RECORD - see doi:10.1042/BJ20081169

THIS IS NOT THE VERSION OF RECORD - see doi:10.1042/BJ20081169

INTRODUCTION

Calcium ions are implicated in the regulation of a wide variety of cellular processes including signal transduction in bacteria, while cytosolic Ca^{2+} levels are tightly controlled by mechanisms involving plasma membrane transporters [1-4]. In contrast to eukaryotic cells, the role of prokaryotic Ca^{2+} transients is less well understood due to the poorly defined regulation of Ca^{2+} influx and efflux systems [3].

In *Escherichia coli*, Ca^{2^+} influx arrangements include the non-proteinaceous complexes of short-chain poly-(R)-3-hydroxybutyrate-polyphopshate- Ca^{2^+} [5, 6]. Complexed poly-(R)-3-hydroxybutyrate (cPHB) is widely found in both prokaryotes and eukaryotes [7] and possesses a number of physiological roles in *E. coli* [6, 8, 9]. Its biosynthesis is unresolved in *E. coli* whilst a periplasmic protein, YdcS, has been reported to exhibit PHB synthase activity only very recently [10]. The non-proteinaceous complexes act as voltage-gated Ca^{2^+} channels with complex gating kinetics, their activity being implicated in the phase of growth-dependent control of cytosolic Ca^{2^+} levels [6, 11, 12]. Moreover, cPHB channels share some characteristics with eukaryotic L-type voltage-activated Ca^{2^+} channels, which are targeted selectively by the Ca^{2^+} channel blockers verapamil, diltiazem and nifedipine, despite their differences in pharmacological properties [13-15].

The biosynthesis and the intracellular distribution of cPHB are regulated by the AtoS-AtoC two-component system (TCS) via its direct action on *atoDAEB* operon transcription upon acetoacetate- or spermidine-mediated induction [16, 17]. In addition, AtoS-AtoC TCS-mediated regulation of cPHB biosynthesis has been reported in the presence of the biogenic amine histamine and the mast cell degranulating synthetic polyamine compound 48/80 (C48/80) [18], which has been shown to induce rapid Ca²⁺ efflux, thus altering the permeability of the outer membrane in *E. coli* [19].

AtoS-AtoC TCS consists of the AtoS histidine kinase and the AtoC response regulator [20] of the NtrC-NifA family of σ^{54} -RNA polymerase transcriptional activators [21] and regulates the expression of the *atoDAEB* operon [21] that encodes for proteins involved in short-chain fatty acid catabolism [22]. AtoS-AtoC TCS plays a fundamental role in *E. coli* physiology and in addition to its participation in a number of interacting regulatory networks [23-26], it may contribute to Ca²⁺ homeostasis through cPHB regulation [16, 17].

E. coli adapt to a wide range of $[Ca^{2+}]e$, as in fresh or sea water [5] and in the gut, where histamine plays prominent physiological roles in gastric acid secretion and inflammation [27], often through TCS-mediated signal transduction [28]. Interestingly, the involvement of Ca^{2+} in histamine release and in the action of the histamine releaser C48/80 is well documented [29], while a limited number of studies argue for the functional implication of Ca^{2+} in regulating proinflammatory signals and for the cross-talk between Ca^{2+} channel blockers, histamine and bacteria-host interactions [30-32].

Since the cPHB biosynthetic pathway in *E. coli* has not been established and taking into consideration the involvement of the *ato* regulon, this study aimed to investigate the effects of extracellular Ca^{2+} in the AtoS-AtoC TCS-regulated cPHB biosynthesis. The reported data provide evidence for an underlying modulatory role of the AtoS-AtoC TCS along with Ca^{2+} -mediated homeostatic mechanisms, leading to new insights on the interaction of signal transduction systems and extracellular Ca^{2+} levels, with potential implications in the symbiotic behavior of bacteria.

MATERIALS AND METHODS

Materials

Histamine dichloride was obtained from SERVA (Heidelberg, Germany), C48/80 from Sigma Chem. Co. (St. Louis, MO, USA) and verapamil hydrochloride from Knoll AG (Ludwigshafen, Germany). All chemical reagents were of analytical grade.

Bacterial strains, plasmids and growth conditions

E. coli K-12 strains BW25113 ($atoSC^+$) [33] and BW28878 ($\Delta atoSC$) [23] were a gift from Hirofumi Aiba (Nagoya University, Japan). Plasmid pUC-Az, containing the *atoS*, *atoC* genes and a part of the *atoDAEB* operon [34], as well as plasmids pEMZ-patoD1 (*atoDAEB* URR-*lacZ* in pDP8), pCPG5 (*atoDAEB* URR-*atoA-lacZ*), pCPG4 (*atoS URR-lacZ*) and pCPG6 (*atoC* URR-*lacZ*) were constructed as described previously [20, 35].

E. coli were grown in LB or in modified M9 mineral medium [16, 18]. AtoS-AtoC TCS was induced by the addition of 10 mM acetoacetate. CaCl₂ or other compounds were added in the culture medium when A_{600} reached 0.3, at concentrations that did not affect cell growth. Cell culture samples were collected at the indicated time points and processed for crotonic acid/cPHB determination.

β-Galactosidase assay

 β -Galactosidase assay was performed using *E. coli* cells carrying the appropriate plasmids [36] and grown in modified M9 mineral medium [16] in the presence of 10 mM acetoacetate or 2.5 mM CaCl₂.

Cell fractionation

E. coli cells were harvested, washed twice with PBS and suspended in 50 mM Tris–HCl, pH 7.5, containing 1 mM phenylmethylsulfonyl fluoride. The cell suspensions were incubated with lysozyme (1 mg/ml) for 30 min at 4 °C and subsequently lysed by sonication. DNase I (20 units/ml) and RNase A (20 µg/ml) were added and the lysates were incubated for 30 min at 4 °C prior to centrifugation at 10,000×g for 20 min at 4 °C. The supernatants were centrifuged at 140,000×g for 1 h at 4 °C to pellet the membranes and separate them from the cytosolic fraction. The membranes were treated in TGS buffer [50 mM Tris–HCl pH 7.5, 10% (v/v) glycerol and 250 mM sucrose] at 4 °C for 1 h and recovered by centrifugation at 170,000×g for 30 min [37].

Determination of cPHB

cPHB determination was carried out by the method of Karr *et al.* [38] as modified by Huang *et al.* [39], based on the acid-catalyzed β -elimination of cPHB to crotonic acid. Dried *E. coli* cell pellets or cytosolic fractions derived from a 10 ml culture sample were incubated with 0.5 ml concentrated H₂SO₄ at 92 °C for 8 h [16, 17]. The samples were placed on ice, 1 ml of saturated sodium sulfate was added and the solution was extracted 4 times with 3 ml of dichloromethane. Sodium hydroxide (100 µl of 1 N) was added to the dichloromethane extracts to convert crotonic acid to sodium crotonate and the dichloromethane was evaporated. The residual was acidified to pH 2 with 3 N sulfuric acid and filtered. The final filtrate was analyzed by HPLC on an Aminex

HPX-87H ion exclusion organic acid analysis column (Bio-Rad), using a Shimadzu (Tokyo, Japan) HPLC chromatography system. Quantitation was performed following comparison of peak absorbance with that of crotonic acid standards (Fluka Buchs, Switzerland). The cPHB content of each sample was determined from the amount of crotonic acid using the established conversion rate [40].

Extraction of cPHB-channels from E. coli cells

E. coli cultures were pelleted by centrifugation and dehydrated by two sequential washes with methanol, 1:1 methanol-acetone and ortacetone [41]. The dehydrated *E. coli* cells or the lyophylized membrane fractions were extracted by incubation in 5 ml chloroform at 4 °C for 36 h. Choloroform was subsequently evaporated with a stream of nitrogen and the amounts of cPHB forming channels were determined as described above, with the exception that the samples were incubated with concentrated H_2SO_4 at 92 °C for 20 min [41].

RESULTS

Involvement of extracellular Ca²⁺ in AtoS-AtoC TCS-mediated cPHB biosynthesis

Exposure of BW25113 (*atoSC*⁺) or BW28878 ($\Delta atoSC$) *E. coli* cells to various [Ca²⁺]e affected cPHB biosynthesis in a concentration-dependent manner, maximal increases being obtained with 2.5 mM and 1 mM, respectively at 0.5 h after extracellular Ca²⁺ addition (Figure 1). The patterns of cPHB levels in the cytosol and membranes were comparable during *E. coli* growth in both strains, with only about 10% being obtained in the membrane fractions compared to the amounts present in the cytosol (Figure 2).

This AtoS-AtoC TCS-mediated regulation of cPHB biosynthesis can be attributed to $[Ca^{2+}]e$, since exposure of cells of both genotypes to equimolar concentrations of EGTA and Ca^{2+} failed to induce any increase in cPHB levels (Figure 3). Moreover, under higher $[Ca^{2+}]e$ and in the presence of MgCl₂, 1 mM EGTA was insufficient to reverse the Ca²⁺-induced increases in cPHB levels (Figure 3).

Effect of extracellular Ca²⁺ on the *ato* operon

Addition of extracellular Ca²⁺ exerted no effect on the minimal *atoDAEB* URR in either *atoSC*⁺ or $\Delta atoSC$ cells, even when they carried the multicopy pUC-Az plasmid (Figure 4). However, it induced *atoDAEB* transcriptional activation acting on an extensive region containing *atoDAEB* URR, *atoD* and a part of *atoA* genes in *atoSC*⁺ cells transformed with plasmid pCPG5 (Figure 4). Furthermore, extracellular Ca²⁺ induced *atoC* transcription in BW25113/pCPG6 cells but exerted no effect on *atoS* transcription in pCPG4-transformed bacteria of both *atoSC*⁺ and $\Delta atoSC$ genotypes (Figure 4).

Effect of verapamil on cPHB biosynthesis

Verapamil elicited an inhibitory effect on cPHB biosynthesis in both $\Delta atoSC$ and AtoS-AtoC overproducing cells, cPHB accumulation decreasing dramatically in the membrane fractions of both strains (Figure 5). Addition of $[Ca^{2+}]e$ increased cPHB levels in verapamil-treated cells. Interestingly, this phenotype was observed in the membranes of $atoSC^+$ but not of $\Delta atoSC$ cells

(Figure 5). Extacellular Ca^{2+} was also able to reverse the diltiazem-mediated cPHB decreases in cells of both genetic backgrounds, this phenotype being circumvented with higher $[Ca^{2+}]e$ in $\Delta atoSC$ cells compared to $atoSC^+$ bacteria (data not shown). Nifedipine caused a more pronounced cPHB reduction that was not reversed by $[Ca^{2+}]e$ in cells of both genotypes (data not shown).

Effect of [Ca²⁺]e on histamine- and C48/80-regulated cPHB biosynthesis

In *atoSC* overexpressing *E. coli*, addition of histamine or C48/80 enhanced the induction of cPHB biosynthesis by 1 mM Ca²⁺, in contrast to the $\Delta atoSC$ cells (Table 1). However, both agents reduced the inductive action of 2.5 mM Ca²⁺ in the $atoSC^+$ strain and induced cPHB production in $\Delta atoSC$ cells (Table 1). At higher [Ca²⁺]e of 5 mM, histamine elicited a minor inductive effect in $atoSC^+$ but not in $\Delta atoSC$ cells (Table 1).

DISCUSSION

Growth of *E. coli* in the presence of increased $[Ca^{2+}]e$ resulted in concentration-dependent induction of cPHB biosynthesis, in both cytosolic and membrane fractions, under an apparent control of the AtoS-AtoC TCS. Maximal cPHB levels accumulated at higher $[Ca^{2+}]e$ in *E. coli* cells overproducing the AtoS-AtoC TCS than in their $\Delta atoSC$ counterparts, which is indicative of a threshold in the Ca²⁺ action in the absence of *atoSC* genes. The specificity of the response was evident by its reversal by the chelating agent EGTA and the restored inductive phenotype upon higher $[Ca^{2+}]e$ and Mg²⁺ addition.

E. coli has no known significant internal Ca²⁺ sequestration machinery, similar for instance to the eukaryotic endoplasmic reticulum. However, Ca²⁺ gradients exist and cPHB seems to control Ca²⁺ storage in *E. coli* cytosol and membranes [5, 6, 42]. Its redistribution from mostly cytosolic to membrane-bound, upon AtoS-AtoC TCS induction, particularly during the stationary phase of growth [17] may have substantial implications in its channel function. Previous reports have implicated cPHB as a possible target of the EGTA-induced rapid decrease in free intracellular Ca²⁺ levels [5], which may be associated with the inhibitory action of EGTA on cPHB biosynthesis in both *atoSC⁺* and *AatoSC* cells. Moreover, although verapamil may elicit effects additional to its putative Ca²⁺-transporter actions [13], the notable decrease in cPHB accumulation in the membrane fractions of both strains, which was reversed by Ca²⁺ addition only in the *atoSC⁺* cells, signified that AtoS-AtoC TCS may regulate predominantly channel cPHB. The significance of the differential effects of the Ca²⁺-channel blockers in the regulatory action of the AtoS-AtoC TCS on cPHB biosynthesis is under investigation.

cPHB induction has been attributed to the *atoDAEB* transcriptional activation by the AtoS-AtoC TCS [16, 17]. Activation in the presence of extracellular Ca^{2+} was observed when an extensive region of the *atoDAEB* sequence was introduced in *E. coli*. Furthermore, the effects of Ca^{2+} on *atoC* but not on *atoS* expression in *atoSC*⁺ cells pointed to different or supplementary mechanisms underlying the action of Ca^{2+} in AtoS-AtoC signaling towards its downstream targets.

The previously reported histamine-induced down-regulation of cPHB biosynthesis in AtoS-AtoC TCS expressing bacteria [18] and the interplay of this biogenic amine with Ca^{2+} in eliciting most of its physiological functions may have immense physiological significance in host-bacteria symbiotic behavior, particularly in the gastrointestinal tract [27-29]. The [Ca²⁺]e-dependent

induction of cPHB biosynthesis by histamine and the functionally related polyamine C48/80 in AtoS-AtoC overproducing but not in $\Delta atoSC$ cells implied the participation of the AtoS-AtoC TCS. These results provide initial indications for a possible Ca²⁺-dependent action of histamine in essential processes of bacterial cells, such as cPHB production, and urge the exploration of yet undefined effects of this host-derived component on the symbiotic *E. coli* through Ca²⁺ transients, beyond the commonly investigated histamine receptor-mediated effects on host (patho)physiology.

In conclusion, several lines of evidence indicate that Ca^{2+} is a key mediator in the regulation of bacterial processes such as motility and chemotaxis [31], as well as in TCS induction [43], establishing its homeostatic role in bacterial physiology and possibly pathophysiology or bacteria-host communication. This first report on the interaction of AtoS-AtoC and Ca²⁺ that govern cPHB biosynthesis provide the lead for the investigation of related signaling networks and their potential exploitation in the treatment of bacterial diseases.

REFERENCES

- 1 Norris, V., Grant, S., Freestone, P., Canvin, J., Sheikh, F.N., Toth, I., Trinei, M., Modha, K. and Norman R. I. (1996) Calcium signaling in bacteria. J. Bacteriol. **178**, 3677-3682
- 2 Holland, I. B., Jones, H. E., Campbell, A. K. and Jacq. A. (1999) An assessment of the role of intracellular free Ca²⁺ in *E. coli*. Biochimie **81**, 901-907
- 3 Dominguez, D.C. (2004) Calcium signaling in bacteria. Mol. Microbiol. 54, 291-297
- Hoffer, S. M., Schoondermark, P., van Veen, H. W. and Tommassen, J. (2001) Activation of gene amplification of pitB, encoding a third phosphate transporter of *Escherichia coli* K-12. J. Bacteriol. 183, 4659-4663
- 5 Jones, H. E., Holland, I. B., Baker, H. L. and Campbell, A. K. (1999) Slow changes in cytosolic free Ca²⁺ in *Escherichia coli* highlight two putative influx mechanisms in response to changes in extracellular calcium. Cell Calcium **25**, 265-274
- 6 Reusch, R. N., Huang, R. and Bramble, L. L. (1995) Poly-3-hydroxybutyrate/ polyphosphate complexes form voltage-activated Ca²⁺ channels in the plasma membranes of *Escherichia coli*. Biophys. J. **69**, 754-766
- 7 Seebach, D., Brunner, A., Burger, H. M., Schneider, J. and Reusch, R. N. (1994) Isolation and ¹H-NMR spectroscopic identification of poly(3-hydroxybutanoate) from prokaryotic and eukaryotic organisms; determination of the absolute configuration (R) of the monomeric unit 3-hydroxybutanoic acid from *Escherichia coli* and spinach. Eur. J. Biochem. **224**, 317-328
- Reusch, R. N., Hiske, T. W. and Sadoff, H. L. (1986) Poly-β-hydroxybutyrate membrane structure and its relationship to genetic transformability in *Escherichia coli*. J. Bacteriol. 168, 553-562
- 9 Xian, M., Fuerst, M. M., Shabalin, Y. and Reusch, R. N. (2007) Sorting signal of *Escherichia coli* OmpA is modified by oligo-(R)-3-hydroxybutyrate. Biochim. Biophys. Acta 1768, 2660-2666
- 10 Dai, D. and Reusch, R. N. (2008) Poly-3-hydroxybutyrate synthase form the periplasm of *Escherichia coli*. Biochem. Biophys. Res. Commun. **374**, 485-489
- 11 Campbell, A. K., Naseem, R., Holland, I. B., Matthews, S. B. and Wann, K. T. (2007) Methylglyoxal and other carbohydrate metabolites induce lanthanum-sensitive Ca²⁺ transients and inhibit growth in *E. coli*, Arch. Biochem. Biophys. **468**, 107-113

- 12 Campbell, A. K., Naseem, R., Wann, K., Holland, I. B. and Matthews, S. B. (2007) Fermentation product butane 2,3-diol induces Ca²⁺ transients in *E. coli* through activation of lanthanum-sensitive Ca²⁺ channels. Cell Calcium **41**, 97-106
- 13 Andersen, C. L., Holland, I. B. and Jacq, A. (2006) Verapamil, a Ca²⁺ channel inhibitor acts as a local anesthetic and induces the sigma E dependent extra-cytoplasmic stress response in *E. coli*. Biochim. Biophys. Acta **1758**, 1587-1595
- 14 McDonald, T. F., Pelzer, S., Trawtwein, W. and Pelszer, D. J. (1994) Regulation and modulation of calcium channels in cardiac, skeletal, and smooth muscle cells. Physiol. Rev. 74, 365-507
- 15 Ling, V. (1997) Multidrug resistance: molecular mechanisms and clinical relevance. Cancer Chemother. Pharmacol. 40, S3-S8 (Suppl.)
- 16 Theodorou, M. C., Panagiotidis, C. A., Panagiotidis, C. H., Pantazaki, A. A. and Kyriakidis, D. A. (2006) Involvement of the AtoS-AtoC signal transduction system in poly-(R)-3-hydroxybutyrate biosynthesis in *Escherichia coli*. Biochim. Biophys. Acta 1760, 896-906
- 17 Theodorou, M. C., Theodorou, E. C., Panagiotidis, C. A. and Kyriakidis, D. A. (2007) Spermidine triggering effect to the signal transduction through the AtoS-AtoC/Az two component system in *Escherichia coli*. Biochim. Biophys. Acta **1770**, 1104-1114
- 18 Kyriakidis, D. A., Theodorou, M. C., Filippou, P. S., Kyriakidis, K. D. and Tiligada, E. (2008) Effect of histamine on the signal transduction of the AtoS-AtoC two component system and involvement in poly-(R)-3-hydroxybutyrate biosynthesis in *Escherichia coli*, Amino acids 35, 45-52
- 19 Katsu, T., Yoshimura, S. and Fujita, Y. (1984) Increases in permeability of *E. coli* outer membrane induced by polycations, FEBS Lett **166**, 175-178
- 20 Lioliou, E. E., Mimitou, E. P., Grigoroudis, A. I., Panagiotidis, C. H., Panagiotidis, C. A. and Kyriakidis, D. A. (2005) Phosphorylation activity of the response regulator of the two-component signal transduction system AtoS-AtoC/Az in *E. coli*, Biochim. Biophys. Acta 1725, 257-268
- 21 Stock, A. M., Robinson, V. L. and Goudreau, P. N. (2000) Two-component signal transduction, Annu. Rev. Biochem. 69, 183-215
- 22 Clark, D. P. and Cronan, Jr. J. E. (1996) Two-carbon compounds and fatty acids as carbon sources. In *Escherichia coli* and *Salmonella typhimurium* (Cellular and Molecular Biology) (Neidhardt, F. C., ed.), vol 1. pp 343-357, American Society of Microbiology, Washington, D.C.
- 23 Oshima, T., Aiba, H., Masuda, Y., Kanaya, S., Masahito, S., Wanner, B. L., Mori, H. and Mizuno, T. (2002) Transcriptome analysis of all two component regulatory system mutants of *Escherichia coli* K-12. Mol. Microbiol. 46, 281-291
- 24 Zhou, L., Lei, X. H., Bochner, B. R. and Wanner, B. L. (2003) Phenotype microarray analysis of *Escherichia coli* K-12 mutants with deletions of all two-component systems. J. Bacteriol. 185, 4956-4972
- 25 Yamamoto, K., Hirao, K., Oshima, T., Aiba, H., Utsumi, R. and Ishihama, A. (2005) Functional characterization *in vitro* of all two-component signal transduction systems from *Escherichia coli*, J. Biol. Chem. **280**, 1448-1456
- 26 Arifuzzaman, M., Maeda, M., Itoh, A., Nishikata, K., Takita, C., Saito, R., Ara, T., Nakahigashi, K., Huang, H.-C., Hirai, A., Tsuzuki, K., Nakamura, S., Altaf-Ul-Amin, M., Oshima, T., Baba, T., Yamamoto, N., Kawamura, T., Ioka-Nakamichi, T., Kitagawa, M.,

THIS IS NOT THE VERSION OF RECORD - see doi:10.1042/BJ20081169

Tomita, M., Kanaya, S., Wada, C. and Mori, H. (2006) Large-scale identification of protein–protein interaction of *Escherichia coli* K-12. Genome Res. **16**, 686-691

- 27 Guarner, F. and Malagelada, J. R. (2003) Gut flora in health and disease. Lancet **361**, 512-519
- 28 Hoch, J. A., Silhavy, T. J. (1995) In Two-component signal transduction (Hoch, J. A. and Silhavy, T. J., eds.), American Society for Microbiology, Washington D.C.
- 29 Chang, W. C. (2006) Store-operated calcium channels and pro-inflammatory signals. Acta Pharmacol Sin. **27**, 813-20
- 30 Hung, C. R. (2007) Protective effect of verapamil on gastric hemorrhagic ulcers in severe atherosclerotic rats. Chin J Physiol. **50**, 315-25
- 31 Tisa, L. S., Sekelsky, J. J. and Adler, J. (2000) Effect of organic antagonists of Ca²⁺, Na⁺ and K⁺ on chemotaxis and motility of *Escherichia coli*, J. Bacteriol. **182**, 4856-4861
- 32 Jones, H. E., Holland, B., Jacq, A., Wall, T. and Campbell, A. K. (2003) *Escherichia coli* lacking the AcrAB multidrug efflux pump also lacks nonproteinaceous, PHB-polyphosphate Ca²⁺ channels in the membrane. Biochim. Biophys. Acta **1612**, 90-97
- Haldimann, A.and Wanner, B. L. (2001) Conditional-replication, integration, excision and retrieval plasmid-host systems for gene structure-function studies of bacteria. J. Bacteriol. 183, 6384-6393
- 34 Canellakis, E. S., Paterakis, A. A., Huang, S. C., Panagiotidis, C. A. and Kyriakidis, D. A. (1993) Identification, cloning, and nucleotide sequencing of the ornithine decarboxylase antizyme gene of *Escherichia coli*. Proc. Natl. Acad. Sci. U.S.A. **90**, 7129-7133
- 35 Matta, M. K., Lioliou, E. E., Panagiotidis, C. H., Kyriakidis, D. A. and Panagiotidis, C. A. (2007) Interactions of the antizyme AtoC with regulatory elements of the *Escherichia coli atoDAEB* operon. J. Bacteriol. **189**, 6324-6332
- 36 Miller, J. H. (1992) In, A short course in bacterial genetics, Cold Spring Harbor Laboratory, Cold Spring Harbor, N.Y.
- 37 Tokishita, S. -I., Yamada, H., Aiba, H. and Mizuno, T. (1990) Transmembrane signal transduction and osmoregulation in *Escherichia coli*: II. The osmotic sensor, EnvZ, located in the isolated cytoplasmic membrane displays its phosphorylation and dephoshorylation abilities as to the activator protein, OmpR. J. Biochem. **108**, 488-493
- 38 Karr, D. B., Waters, J. K. and Emerich, D. W. (1983) Analysis of poly-β- hydroxybutyrate in Rhizobium japonicum bacteroids by ion-exclusion high-pressure liquid chromatography and UV detection. Appl. Environ. Microbiol. **46**, 1339-1344
- 39 Huang, R. and Reusch, R. N. (1996) Poly(3-hydroxybutyrate) is associated with specific proteins in the cytoplasm and membranes of *Escherichia coli*. J. Biol. Chem. **271**, 22196-22202
- 40 Tagushi, S., Maehara, A., Takase, K., Nakahara, M., Nakamura, H. and Doi, Y. (2001) Analysis of mutational effects of a polyhydroxybutyrate (PHB) polymerase on bacterial PHB accumulation using an in vivo assay system, FEMS Microbiol. Lett. **198**, 65-71
- 41 Pavlov, E., Grimbly, C., Diao, C. T. and French, R. J. (2005) A high-conductance mode of a poly-3-hydroxybutyrate/calcium/polyphosphate channel isolated from competent *Escherichia coli* cells, FEBS Lett **579**, 5187-5192
- 42 Michiels, J., Xi, C., Verhaert, J. and Vanderleyden, J. (2002) The functions of Ca²⁺ in bacteria: a role for EF-hand proteins? Trends Microbiol. 10, 87-93
- 43 Rampersaud, A., Utsumi, R., Delgado, J., Forst, S. A. and Inouye, M. (1991) Ca²⁺enhanced phopshorylation of a chimeric protein kinase involved with bacterial signal transduction. J. Biol. Chem. **266**, 7633-7637

Figure legends

Figure 1 Dose-response of extracellular Ca²⁺ on cPHB biosynthesis

Total cPHB levels in BW25113/pUC-Az and BW28878 *E. coli* cells at 0.5 h after addition of 0.25-5 mM CaCl₂ in the culture medium. Results are presented as the average of two experiments. The error for each data point is less than 5 %.

Figure 2 Effect of extracellular Ca²⁺ on cPHB levels during *E. coli* growth

Total cPHB levels in the cytosolic (closed symbols) and membrane (open symbols) fractions of (A) BW25113/pUC-Az and (B) BW28878 cells grown in the absence (\blacksquare , \Box) or presence of 1 mM (\bullet , \circ) or 2.5 mM (\bullet , \diamond) CaCl₂. Results are presented as the average of two experiments. The error for each data point is less than 5 %.

Figure 3 Effect of EGTA and MgCl₂ on Ca²⁺-induced cPHB biosynthesis

(A) BW25113/pUC-Az and (B) BW28878 *E. coli* cells were grown in the absence (\blacksquare) or presence of 1 mM EGTA, either alone (\blacktriangle) or in combination with 1 mM (\bullet , \circ) or 2.5 mM (\bullet , \diamond) CaCl₂, without (closed symbols) or with (open symbols) the addition of 1 mM MgCl₂. Results are presented as the average of two experiments. The error for each data point is less than 5 %.

Figure 4 Effect of Ca²⁺ on *atoDAEB*, *atoC* and *atoS* transcription

Activation of *atoDAEB*, *atoC* and *atoS* promoters in (A) BW25113 and (B) BW28878 *E. coli* cells transformed with plasmids *atoDAEB* URR-*lacZ* without (pEMZ-patoD1) or with pUC-Az (pUC-Az/pEMZ-patoD1), *atoDAEB* URR-*atoD-atoA-lacZ* (pCPG5), *atoC* URR-*lacZ* (pCPG6) and *atoS* URR-*lacZ* (pCPG4), in the absence (open columns) or presence of 10 mM acetoacetate (shaded columns) or 2.5 mM CaCl₂ (filled colums). Results are presented as the average of two experiments. The error for each data point is less than 5 %.

Figure 5 Effect of verapamil on AtoS-AtoC regulated cPHB biosynthesis

Total cPHB levels in the cytosolic (closed symbols) and membrane (open symbols) fractions of (A) BW25113/pUC-Az and (B) BW28878 cells grown in the absence (\blacksquare , \Box) or presence of 500 μ M verapamil alone (\bullet , \circ) or with 5 mM CaCl₂ (\bullet , \diamond). Results are presented as the average of two experiments. The error for each data point is less than 5 %.

B

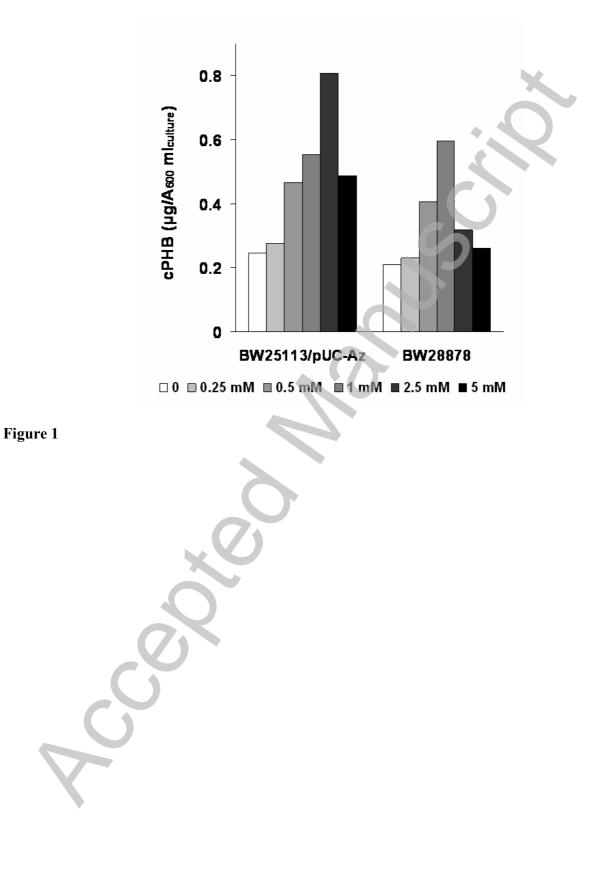
Iournal Immediate Publication. Published on 14 Oct 2008 as manuscript BJ20081169

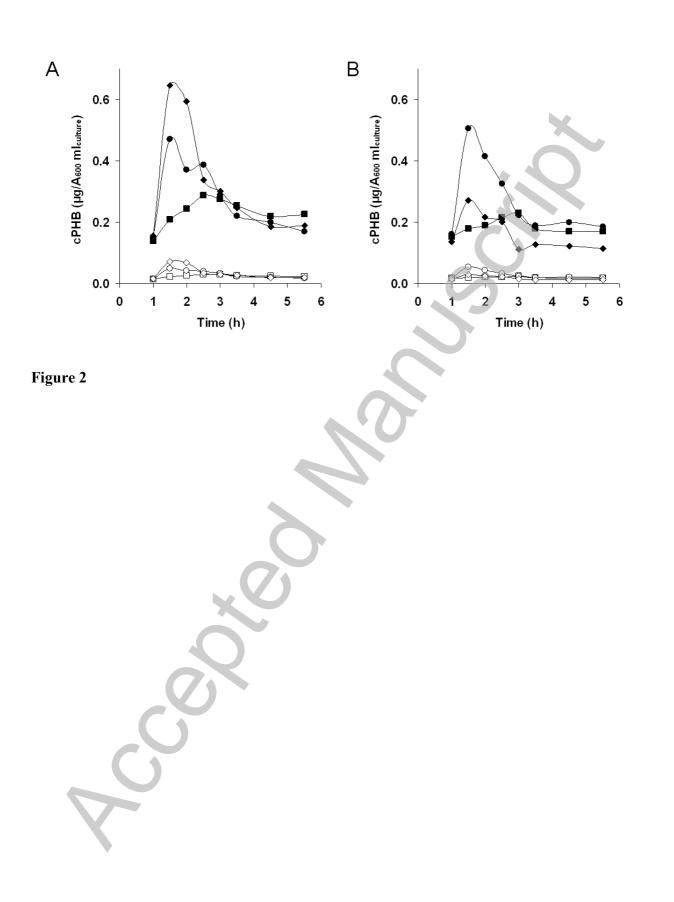
The results are expressed as $\mu g \ cPHB/A_{600} \ ml_{culture}$ and presented as the means $\pm S.D.$ of two experiments in *E. coli* BW25113/pUC-Az and BW28878.

3 11 2 0070.					<u> </u>	
		BW25113/pUC-Az			BW28878	()
CaCl ₂		+ Histamine	+ C48/80		+ Histamine 85 mM	+ C48/80
mM		5 mM	10 µg/ml		Ö5 mM	10 μg/ml
0	0.24 ± 0.01	0.16 ± 0.01	0.39 ± 0.01	0.21 ± 0.01	0.043 ± 0.01	0.12 ± 0.01
1	0.55 ± 0.03	0.65 ± 0.03	0.84 ± 0.04	0.60 ± 0.02	$0 \frac{1}{2} 7 \pm 0.02$	0.51 ± 0.02
2.5	0.81 ± 0.04	0.55 ± 0.02	0.23 ± 0.01	0.20 ± 0.01	0 = 0.01	0.32 ± 0.01
5	0.39 ± 0.02	0.47 ± 0.02	0.22 ± 0.01	0.28 ± 0.01	$0 \frac{2}{9} \pm 0.01$	0.31 ± 0.01
P	cce				THIS IS NOT THE	

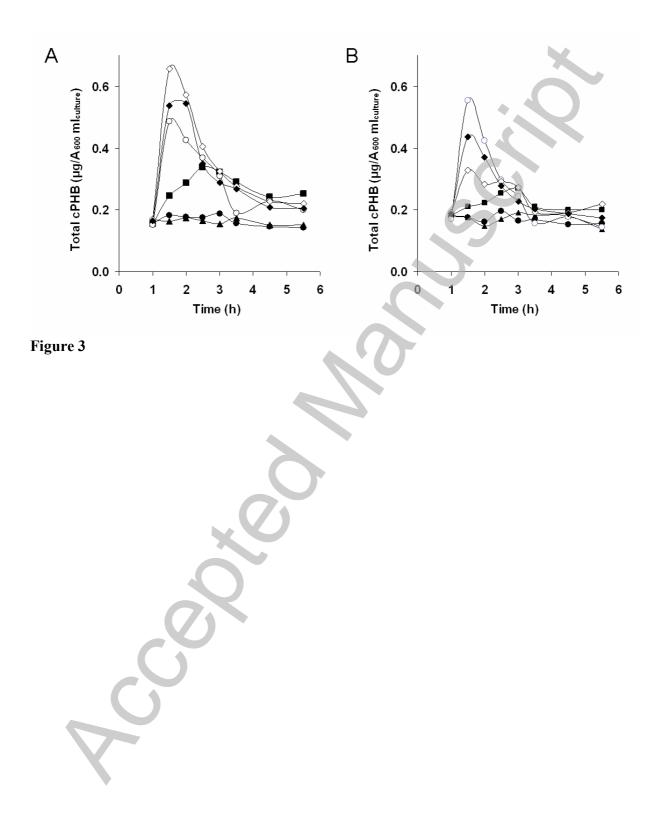
Licenced copy. Copying is not permitted, except with prior permission and as allowed by law. © 2008 The Authors Journal compilation © 2008 Biochemical Society

B

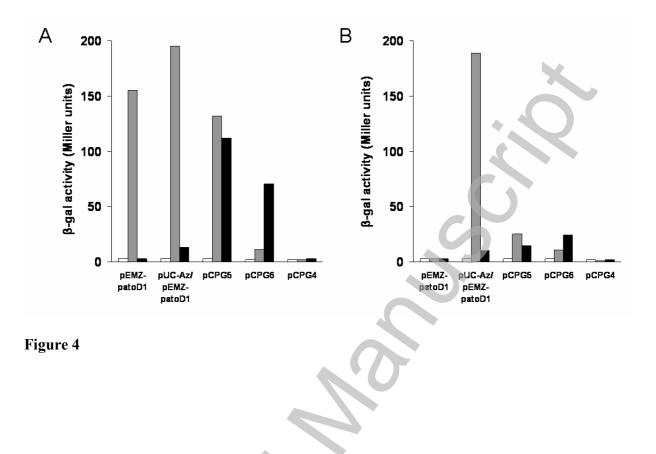




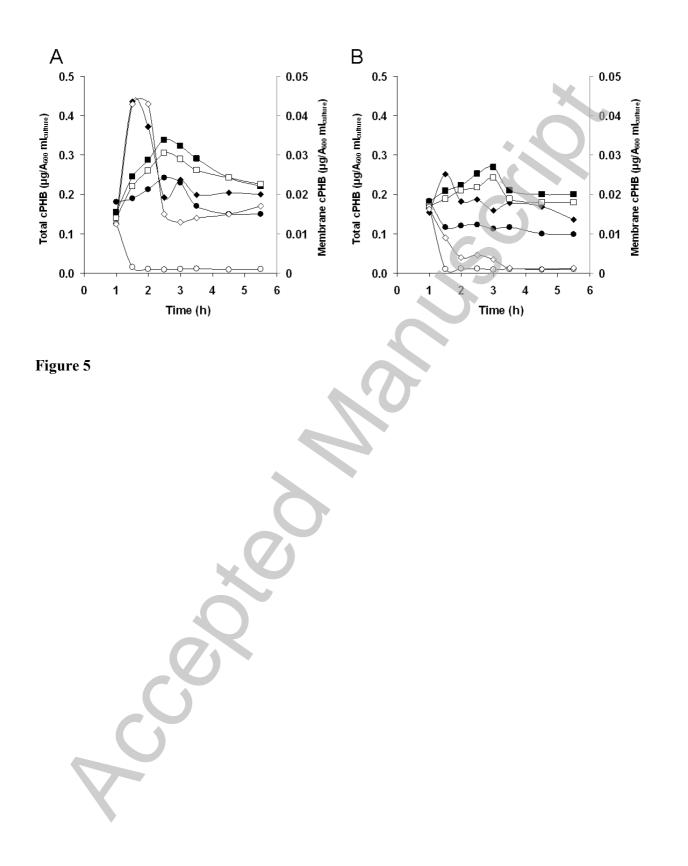
BJ



BJ



B



16