2007

pH of the Cytoplasm and Periplasm of Escherichia coli: Rapid Measurement by Green Fluorescent Protein Fluorimetry

Joan Slonczewski
Kenyon College, slonczewski@kenyon.edu

Follow this and additional works at: http://digital.kenyon.edu/biochem_publications
Part of the Biochemistry, Biophysics, and Structural Biology Commons, and the Biology Commons

Recommended Citation
pH of the Cytoplasm and Periplasm of *Escherichia coli*: Rapid Measurement by Green Fluorescent Protein Fluorimetry

Jessica C. Wilks and Joan L. Slonczewski*

Department of Biology, Kenyon College, Gambier, Ohio 43022

Received 20 April 2007/Accepted 21 May 2007

Cytoplasmic pH and periplasmic pH of *Escherichia coli* cells in suspension were observed with 4 s time resolution using fluorimetry of TorA-green fluorescent protein mutant 3* (TorA-GFPmut3*) and TetR-yellow fluorescent protein. Fluorescence intensity was correlated with pH using cell suspensions containing 20 mM benzoate, which equalizes the cytoplasmic pH with the external pH. When the external pH was lowered from pH 7.5 to 5.5, the cytoplasmic pH fell within 10 to 20 s to pH 5.6 to 6.5. Rapid recovery occurred until about 30 s after HCl addition and was followed by slower recovery over the next 5 min. As a control, KCl addition had no effect on fluorescence. In the presence of 5 to 10 mM acetate or benzoate, recovery from external acidification was diminished. Addition of benzoate at pH 7.0 resulted in cytoplasmic acidification with only slow recovery. Periplasmic pH was observed using TorA-GFPmut3* exported to the periplasm through the Tat system. The periplasmic location of the fusion protein was confirmed by the observation that osmotic shock greatly decreased the periplasmic fluorescence signal by loss of the protein but had no effect on the fluorescence of the cytoplasmic protein. Based on GFPmut3* fluorescence, the pH of the periplasm equaled the external pH under all conditions tested, including rapid acid shift. Benzoate addition had no effect on periplasmic pH. The cytoplasmic pH of *E. coli* was measured with 4 s time resolution using a method that can be applied to any strain construct, and the periplasmic pH was measured directly for the first time.

In order to colonize the human gastrointestinal tract, the enteric bacterium *Escherichia coli* must be able to grow between pH 4.5 and pH 9 (7). Over this wide pH range, *E. coli* preserves enzyme activity, as well as protein and nucleic acid stability, by maintaining the cytoplasmic pH in the range from pH 7.2 to 7.8 (26, 27, 32). *E. coli* responds rapidly to intracellular pH change; after acidification of the external environment, the intracellular pH of *E. coli* begins to recover within 1 min, and full recovery occurs within 5 min (28). The efficiency with which *E. coli* maintains pH homeostasis has been attributed to a combination of constitutive and regulated mechanisms, but the essential requirements remain poorly understood (7, 9, 14, 18, 28). Some components of pH homeostasis act in the presence of chloramphenicol, whereas others require ongoing protein synthesis (10).

Previously, cytoplasmic pH has been measured using 31P nuclear magnetic resonance (NMR) of titratable phosphate and methylphosphonate (28) and through transmembrane equilibration of radiolabeled permeant acids (32). Both methods have limitations. Radiolabeled permeant acids have low resolution using fluorimetry of plasmid-expressed GFP mutant 3* (GFPmut3*) and YFP gene fusions. The GFPmut3* fusion strain was expressed with a TorA signal peptide that can either retain the fusion protein in the cytoplasm or direct its transport to the periplasm (3, 17, 30). To our knowledge, there has been no previous report of measurement of periplasmic pH.

In order to measure the pH of *E. coli*, we devised a procedure based on fluorimetry of plasmid-expressed GFP mutant 3* (GFPmut3*) and YFP gene fusions. The GFPmut3* fusion strain was expressed with a TorA signal peptide that can either retain the fusion protein in the cytoplasm or direct its transport to the periplasm (3, 17, 30).

In previous reports, the pH-dependent changes in fluorescence have been observed using fluorescence microscopy (22). Microscopy, however, necessitates cumbersome quantitation procedures that introduce error and limit the time scale of the observable signal. An alternative approach is fluorescence spectroscopy or fluorimetry, a sensitive technique that allows observation of a live cell culture in liquid medium. Fluorimetry has not yet been widely used to measure changes in the intracellular pH of cell suspensions (1), in part because of the need for highly sensitive instrumentation which has only recently become available.

In order to measure the pH of *E. coli*, we devised a procedure based on fluorimetry of plasmid-expressed GFP mutant 3* (GFPmut3*) and YFP gene fusions. The GFPmut3* fusion strain was expressed with a TorA signal peptide that can either retain the fusion protein in the cytoplasm or direct its transport to the periplasm (3, 17, 30). To our knowledge, there has been no previous report of measurement of periplasmic pH.

**Materials and Methods**

**Strains and growth conditions.** All strains were derivatives of *E. coli* K-12. Strain W3110 was transformed with pSL38-YFP (12) to form strain JLS0617. Strains MC4100AR ΔluxABCDE/pBAD TorA-GFPmut3* and MC4100/pBAD TorA-GFPmut3* were obtained from Mullineaux et al. (17).
Strain JLS0617 was cultured overnight in buffered M63 salts medium [20 mM homopiperazine-N,N′-bis-2-(ethanesulfonic acid) (HOMOPIPES), pH 7.5] containing 1.5% casein hydrolysate, 0.8% glycerol, and 100 µg/ml ampicillin. The overnight culture was diluted 25-fold in the same medium and cultured to late log phase (OD600 0.8 to 0.9) at 37°C with aeration.

The MC4100 strains containing pAra TorA-GFPmut3* were grown overnight in potassium-modified Luria broth with NaCl replaced by 100 mM KCl (designated LBK medium) as described by Maurer et al. (14). The LBK medium was buffered with 20 mM HOMOPIPES, pH 7.5. To maintain the plasmid, media included 50 µg/ml ampicillin. Each overnight culture was diluted 1,000-fold into the same medium with 200 µM L-arabinose. Bacteria were cultured to late log phase (OD600 0.8 to 0.9) at 37°C with aeration.

For cytoplasmic pH measurement, the cultures were resuspended to an OD600 of 0.4 in M63 medium containing 1.5% casein hydrolysate, 0.8% glycerol, and HOMOPIPES (5 to 50 mM depending on the experiment). The pH was adjusted with KOH to pH 5.5 to 8.0, depending on the experiment. For periplasmic pH measurement, cultures of the MC4100/pAra TorA-GFPmut3* strain (with a functional tat system) were washed and resuspended to an OD600 of 0.5 in 20 ml of buffered LBK medium (20 mM HOMOPIPES, pH 7.5) containing ampicillin (50 µg/ml) and no arabinose. The cultures were incubated at 37°C for a further 3 h to allow complete transport of TorA-GFPmut3* to the periplasm (3). Then cultures were resuspended in M63 medium (supplemented as described above) at an OD600 of 0.4. All resuspended cultures were stored on ice until fluorimetry.

**Fluorescence spectroscopy.** Excitation spectra were recorded using a FluoroMax-3 spectrophotofluorimeter (Horiba Jobin Yvon). This instrument has a high signal-to-noise ratio (Water Raman signal-to-noise ratio, 3,000:1) and an emission detector employing photon counting in order to detect low light levels. A cell suspension (3 ml) was placed into a Starna Spectrosil quartz cuvette with a path length of 10 mm. Aliquots of solution were added through an injection port using a 1-ml syringe. The temperature of the chamber was adjusted to 30°C, and aeration was provided by stirring. GFPmut3* excitation was measured from 480 to 510 nm (slit width, 2 nm), using an emission wavelength of 545 nm (slit width, 20 nm) (6). YFP excitation was measured from 450 to 520 nm (slit width, 2 nm), using an emission wavelength of 550 nm (slit width, 20 nm) (15). Spectra were recorded for three biological replicates at each pH. For time course experiments, continuous excitation spectra were obtained every 4 s for 1 min before aliquot addition (time zero) and then for 5 min after addition.

Data were fitted to a standard curve correlating internal pH with fluorescence intensity. The standard curve was determined for each fluorophore (GFPmut3* and YFP) by obtaining fluorescence measurements of samples resuspended at pH 5.5, 6.0, 7.0, 7.5, and 8.0, with inclusion of 20 mM sodium benzoate, a permeant acid that equilibrates cytoplasmic pH with external pH. For GFP fluorescence, an equation was fitted to the curve of GFPmut3* signal intensity (GFPmut3* signal intensity = −2.2 × 1010 + 5.8 × 108 × pH) and was used to convert time course signal intensities (sum of 480 nm to 510 nm) to pH units. For YFP, the equation was as follows: YFP signal intensity = −3.91 × 109 + 9.4 × 107 × pH. For each fluorophore, the equation was used to fit a standard curve to the data of a given experiment, interpolating between intensities at pH 5.5 and intensities at pH 7.5. The pH curve was modified for experiments in which a permeant acid was added to cultures in medium with an external pH of 7.0. In this case, linear interpolation was conducted between pH 7.6 and pH 7.0. The fluorescence corresponding to pH 7.0 was determined by adding an additional 20 mM benzoate to each test sample at the end of the time course.

**Osmotic shock.** To permeabilize the outer membrane and remove the periplasmic contents, osmotic shock was performed. The cultures were spun down and resuspended in buffered sucrose-EDTA (20% sucrose, 33 mM morpholinepropanesulfonic acid [MOPS] [pH 7.3], 100 µM EDTA) and then were held at room temperature for 10 min. The cultures were resuspended in ice-cold 500 µM MgCl2. Cells were washed and resuspended to an OD600 of 0.4 in buffered (5 mM HOMOPIPES) supplemented M63 medium adjusted to pH 5.5 or to pH 7.5.

**RESULTS**

**pH dependence of GFPmut3** and YFP in the living cell. The fluorescence excitation spectra of cytoplasmic TorA-GFPmut3* (17) and TetR-YFP (12) were observed as a function of cytoplasmic pH (Fig. 1A and B). Previously, GFP has been used as a probe for determination of bacterial pH by fluorescence microscopy but not by fluorimetry of cell suspensions (22). Cells of strains MC4100AR ΔtatABCDE TorA-GFPmut3* and JLS0617 (W3110/pSL38_YFP) were suspended in buffered M63 medium with supplements, adjusted to pH values in the range from pH 5.5 to 8.0. The permeant acid benzoate (20 mM) was included in order to equalize the cytoplasmic and extracellular pHs.

The excitation spectra for cell suspensions containing TorA-GFPmut3* and YFP peaked at excitation wavelengths of approximately 511 nm and 514 nm, respectively (Fig. 1A and B). These values correspond to the peak excitation wavelengths reported for GFPmut3* and YFP in vitro (6, 15). Overall, for both strains, the fluorescence signal increased with increasing pH over the range from pH 5.5 to 8.0. In the pH range relevant to cytoplasmic pH perturbation, typically pH 7.0 to 8.0, the YFP signal showed greater variation than the GFPmut3* signal. The GFPmut3* signal, however, showed fivefold-greater intensity than the YFP signal.

For both fluorophores, the fluorescence signal was found to be critically dependent on cell density. When the cell density was too low (OD600 < 0.3), the fluorescence signal intensity was undetectable. At higher cell densities, however (OD600 > 0.4), the scattering of the excitation or emission beam from the concentrated sample reduced the apparent signal intensity.
For all our fluorescence observations, the cells were resuspended at an OD$_{600}$ of 0.4.

To use the GFPmut3* and YFP signals as a measure of cytoplasmic pH, standard curves were generated for pH dependence. For each excitation spectrum, the fluorescence intensities were summed over a peak wavelength range. The peak fluorescence signal was observed for cultures suspended in medium over a range of pH values in the presence of the permeant acid benzoate (20 mM) in order to peg the cytoplasmic pH to the pH of the medium (Fig. 2). The intensity of the fluorescence signal increased with increasing pH over the range expected for each fluorophore, including the range of interest for *E. coli* growth, pH 5.5 to 8.0. The YFP signal varied more than the GFPmut3* signal over the cytoplasmic range of pH 7.0 to 8.0. The GFPmut3* signal, however, showed nearly 10-fold-greater intensity than the YFP signal; thus, overall the change in intensity over the cytoplasmic range from pH 7.0 to 8.0 was fivefold greater for GFPmut3* than for YFP.

The regression equations generated from these standard curves were used to convert signal intensities (defined as the sum of intensities at 480 nm to 510 nm) to pH units in order to estimate the cytoplasmic pH values during time course experiments. The equations were fitted to the data by interpolating between fluorescence intensities observed in cells with the cytoplasmic pH depressed by 20 mM benzoate so as to equal the extracellular pH, at external values of pH 5.5 and pH 7.5.

**Cytoplasmic pH shift.** A time course analysis was conducted in order to observe cytoplasmic pH and recovery following rapid acidification of the external medium (Fig. 3). Strains MC4100AR ΔtatABCDE TorA-GFPmut3* and JLS0617 were suspended in mildly buffered M63 medium (5 mM HOMOPIPES) adjusted to pH 7.5. At time zero, HCl (8.5 mM) was added in order to lower the external pH to about pH 5.5. After the completion of the time course, the external pHs of the cultures were measured and found to be between pH 5.5 and 5.7.

Before HCl addition, the intracellular pH of all cultures was pH 7.6, as determined based on a standard curve obtained using cells from the same culture. After HCl addition, the cytoplasmic pH of each sample fell within 10 to 20 s to pH 5.6 to 6.5. Recovery began within 4 s after the lowest point (Fig. 3A and B). The extent of internal pH change varied similarly between the strains and between the biological replicates. In both strains (one strain containing GFPmut3* and one strain containing YFP), rapid recovery occurred until about 30 to 60 s after HCl addition and was followed by slower recovery over the next several minutes.

**Cytoplasmic pH shift in the presence of permeant weak acids.** The effect of permeant acid addition on cytoplasmic pH was tested. A permeant weak acid dissociates upon entering the cell, partly equilibrating cytoplasmic and external pHs and partly impairing pH homeostasis; the precise balance is poorly understood and is a complex function of cell buffering capacity and ion transport (25, 26, 28).

The effects of acetate on cell pH are of particular interest because *E. coli* produces acetate rapidly during log-phase aerobic growth on most substrates (29), as well as during ferment-
Cultures of strain MC4100AR ΔtatABCDE TorA-GFPmut3* were suspended in LBK medium weakly buffered at pH 7.5 in the presence of 0, 1, 5, and 20 mM acetate (Fig. 4). At time zero, addition of HCl (8.5 mM) caused a rapid fall in the cytoplasmic pH, followed by a relatively slow partial recovery of the cytoplasmic pH in the presence of acetate. A similar experiment was conducted to observe the time course of cytoplasmic acidification in the presence of 0, 1, 5, and 20 mM benzoate (Fig. 5A). Following HCl addition, the cytoplasmic pH of all cultures fell within 10 to 15 s to the low point. The initial rates of recovery were similar over the range from 0 to 5 mM benzoate. The subsequent slow recovery, however, was strongly dependent upon the benzoate concentration, which ranged from 0 mM to 20 mM. The cytoplasmic pH of the cultures suspended in the absence of benzoate dropped to approximately pH 6.5 and then recovered fully, whereas the internal pH of the cultures containing 20 mM benzoate dropped to pH 5.5 (the pH of the external medium) and showed minimal recovery.

In another experiment, benzoate (5, 10, and 20 mM) or pH 7.0 buffer was added to cultures buffered at pH 7.0 without changing the external pH (Fig. 5B). To calibrate the pH, a final addition of 20 mM benzoate was made, which set the cytoplasmic pH at or near the pH of the external medium (pH 7); then the final fluorescence intensity was used to calibrate pH 7.0. The initial addition of benzoate transiently acidified the cytoplasm (Fig. 5B). No rapid short-term recovery was observed (Fig. 3 and 4A). The rate of long-term recovery was dependent on the benzoate concentration. In the presence of 5 mM benzoate, the cytoplasmic pH fell to pH 7.3, close to the pH of the external medium (pH 7.0), and made nearly a full recovery. The presence of 10 and 20 mM benzoate, however, lowered the cytoplasmic pH (Fig. 5B). In the presence of these higher concentrations of benzoate the cytoplasmic pH exhibited minimal recovery. Changes in the signal unrelated to benzoate were controlled for by the addition of pH 7.0 buffer at time zero (Fig. 5B), which caused no significant change in pH.

Controls for chloride and buffer change. Control experiments were designed to eliminate the possibility of effects on the fluorescence signal unrelated to pH change. The factors tested include chloride ion concentration, which is known to quench the fluorescence signal of some GFP derivatives (31); buffer addition at near-cytoplasmic pH (pH 7.5); and buffer addition at low pH (pH 5.5). Strain MC4100AR ΔtatABCDE TorA-GFPmut3* was suspended in strongly buffered M63 minimal medium adjusted to either pH 5.5 or pH 7.0. At time zero, either KCl (8.5 mM; pH 7.5) or HOMOPIPES (50 mM, pH 7.5) was added instead of HCl. Benzoate (20 mM) was added at 4.1 min for pH calibration, as described in Materials and Methods. For each condition, results from three independent cultures were averaged.
pH dependence of periplasmic GFPmut3*. GFP fusion proteins are incapable of folding within the periplasm; thus, GFP functioning in the periplasm requires a system capable of transporting the fully folded protein into the periplasmic space (8, 30). In bacteria, fully folded proteins are transported into the periplasm using the tat pathway (4, 30). By fusing GFP to a protein normally exported by tat, fully folded active GFP can be transported into the periplasm (3, 17, 30). For our observations of periplasmic pH, we used an MC4100 construct with a functional tat system to transport TorA-GFPmut3* to the periplasm (17).

Strain MC4100AR TorA-GFPmut3* was cultured to late log phase in the presence of L-arabinose. One half of the cell culture was spun down, washed, and suspended in the absence of arabinose. Cultures were suspended in supplemented M63 medium (50 mM HOMOPIPES) at pH 7.5 and at pH 5.5. For each condition, three independent cultures were tested. pH_ex, external pH.

Strain MC4100AR TorA-GFPmut3* was cultured as described in Materials and Methods. Cells were harvested following (A) no further incubation and (B) 3 h of incubation in the absence of arabinose. Cultures were suspended in supplemented M63 medium (50 mM HOMOPIPES) at pH 7.5 and at pH 5.5. For each condition, three independent cultures were tested. pH_ex, external pH.

Osmotic shock of tat and tat+/H11545 strains. In order to demonstrate that GFPmut3* was successfully exported into the periplasm, cells incubated without arabinose were subjected to osmotic shock (Fig. 7). Osmotic shock permeabilizes the outer membrane and causes the periplasmic contents to leak into the extracellular space, while the inner membrane and cytoplasm remain intact (2). Osmotic shock was performed on a tat+/H11001 strain (MC4100AR TorA-GFPmut3*) and on a tat deletion strain (MC4100AR ΔtatABCDE TorA-GFPmut3*) in which the TorA-GFPmut3* remains in the cytoplasm. One half of each culture was subjected to osmotic shock, and one half was left unshocked. All cultures were suspended to the same pH 5.5 or pH 7.5 (Fig. 6A). This observation is consistent with little or no TorA-GFPmut3* transport into the periplasm. After 3 h of incubation without L-arabinose, however, the excitation intensity showed a substantial dependence on the external pH (Fig. 6B). This finding is consistent with the transport of all or most of the TorA-GFPmut3* into the periplasm, whose pH is presumed to remain at or near the pH of the external medium (19).

Osmotic shock of tat and tat+/H11545 strains. In order to demonstrate that GFPmut3* was successfully exported into the periplasm, cells incubated without arabinose were subjected to osmotic shock (Fig. 7). Osmotic shock permeabilizes the outer membrane and causes the periplasmic contents to leak into the extracellular space, while the inner membrane and cytoplasm remain intact (2). Osmotic shock was performed on a tat+/H11001 strain (MC4100AR TorA-GFPmut3*) and on a tat deletion strain (MC4100AR ΔtatABCDE TorA-GFPmut3*) in which the TorA-GFPmut3* remains in the cytoplasm. One half of each culture was subjected to osmotic shock, and one half was left unshocked. All cultures were suspended to the same pH 5.5 or pH 7.5 (Fig. 6A). This observation is consistent with little or no TorA-GFPmut3* transport into the periplasm. After 3 h of incubation without L-arabinose, however, the excitation intensity showed a substantial dependence on the external pH (Fig. 6B). This finding is consistent with the transport of all or most of the TorA-GFPmut3* into the periplasm, whose pH is presumed to remain at or near the pH of the external medium (19).
Fluorescence probes for cytoplasmic and periplasmic pHs.

The reliability of our method was strengthened by the use of two different gene fusion constructs, TorA-GFPmut3* and TetR-YFP. The two probes showed significant differences in pH titration (Fig. 2), and the GFPmut3* fluorophore showed fivefold-greater signal intensity than YFP; these effects were consistent with previous observations of GFPmut3* and YFP (6). Nevertheless, the two probes generated essentially the same profiles of cytoplasmic pH response to an acid shift (Fig. 3).

The Tat-transported TorA-GFPmut3* fusion enabled pH measurement in the periplasm, a cell compartment where pH has not been measured previously (Fig. 8). The direct measurement of periplasmic pH is of interest because the periplasm is the gram-negative cell’s frontline defense against pH change, containing major pH stress proteins such as the HdeA and HdeB acid-dependent periplasmic chaperones (19, 21, 29). These chaperones are needed to respond to protein misfolding during acid stress.

Previously, GFP fusion proteins have not been observed in the periplasm because the protein fails to fold properly after transport and is thus inactive (8). However, the Tat twin-arginine translocase pathway, unlike the Sec pathway, allows the transport of fully folded globular proteins into the periplasm.
(4, 30). Thus, the GFPMut3* exported to the periplasm in folded form via this pathway is as active as the GFPMut3* located within the cytoplasm. Periplasmic transport requires removal of arabinose; in the presence of arabinose, the fusion protein remains in the cytoplasm. Thus, our approach allows measurement of cytoplasmic pH and periplasmic pH in the same Tat strain.

Regulation of cytoplasmic pH after rapid external acid shift. Previous kinetic studies of intracellular pH regulation used 31P NMR of titratable phosphates, a method that requires highly concentrated cell suspensions (27, 28). In these experiments, cytoplasmic pH does not begin to recover until approximately 2 min after HCl addition. The initial dip before recovery lasts significantly longer than the 10 to 20 s that was reported in the present study. A possible reason for this difference may be the relatively stressed condition of the cells in the NMR experiment, which were harvested in late log phase and resuspended at high density.

The conditions of the experiments in the present study provided sufficient resolution to distinguish two phases of cytoplasmic pH recovery: the initial recovery, within 1 min after the dip, followed by a slower recovery over several minutes. We showed that the fast recovery is absent when pH is perturbed by a permeant acid such as benzoate, while partial slow recovery remains. Future experiments will enable us to test the two phases of pH recovery in various mutant backgrounds under different physiological conditions.

ACKNOWLEDGMENTS

This work was supported by grant MCB-0644167 from the National Science Foundation.

We thank Colin Robinson and Melanie Berkmen for the generous gift of strains. We thank Scott Cummings in the Kenyon Chemistry Department for use of the Jobin fluorimeter, which was purchased through an undergraduate education grant to Kenyon College from the Howard Hughes Medical Institute Biomedical Sciences Education Program. We thank Daniel Barich for computing assistance and China Ugwu for technical assistance.

REFERENCES