

Identification of Genes Encoding the Folate- and Thiamine-Binding Membrane Proteins in Firmicutes^{∇‡}

Aymerick Eudes,^{1†} Guus B. Erkens,^{2†} Dirk J. Slotboom,² Dmitry A. Rodionov,^{3,4}
Valeria Naponelli,⁵ and Andrew D. Hanson^{1*}

Horticultural Sciences Department, University of Florida, Gainesville, Florida 32611¹; Department of Biochemistry, University of Groningen, Groningen Biomolecular Science and Biotechnology Institute, Nijenborgh 4, 9747 AG Groningen, The Netherlands²; Burnham Institute for Medical Research, La Jolla, California 92037³; Institute for Information Transmission Problems RAS, Moscow 127994, Russia⁴; and Food Science and Human Nutrition Department, University of Florida, Gainesville, Florida 32611⁵

Received 30 July 2008/Accepted 27 August 2008

Genes encoding high-affinity folate- and thiamine-binding proteins (FolT, ThiT) were identified in the *Lactobacillus casei* genome, expressed in *Lactococcus lactis*, and functionally characterized. Similar genes occur in many Firmicutes, sometimes next to folate or thiamine salvage genes. Most *thiT* genes are preceded by a thiamine riboswitch.

The folate and thiamine transport systems of *Lactobacillus casei* were partially characterized 30 years ago by Henderson and colleagues (8, 9, 11, 12). These systems were shown to involve two small membrane proteins for specific substrate binding—one for folate and the other for thiamine—as well as an uncharacterized component shared by both systems.

To identify genes encoding the binding proteins (FolT and ThiT), we used the AACompIdent tool on the ExpASY server (27) to search the *L. casei* (strain ATCC 334) genome for open reading frames with amino acid compositions and molecular masses matching those published for FolT and ThiT (9, 12). The best match for FolT was LSEI_2252, a 19.0-kDa protein with five predicted transmembrane domains (Fig. 1A). LSEI_2252 has homologs in other Firmicutes, and in some cases, the corresponding genes are adjacent to *folC* (Fig. 1B). FolC is a salvage enzyme that mediates polyglutamylation of folates (2). The best match for ThiT was LSEI_1757, a 21.2-kDa protein with six predicted transmembrane domains, which belongs to the YuaJ family (InterPro accession number IPR012651) of predicted, uncharacterized thiamine transporters in the *Bacillus/Clostridium* group (20). LSEI_1757 is 32% identical to *Bacillus subtilis* YuaJ (Fig. 1C). In several Firmicutes, the *thiT* gene forms a putative operon with the thiamine pyrophosphokinase *thiN* gene (Fig. 1D). Like FolC, ThiN is a salvage enzyme that converts thiamine to its active pyrophosphate form (15).

To investigate whether *folT* and *thiT* indeed code for vitamin-binding proteins, the *folT* and *thiT* genes were PCR amplified from *L. casei* genomic DNA, cloned between the NcoI and SstI sites of pNZ8048, a vector carrying the nisin-inducible *nisA* promoter (14), and introduced into *Lactococcus lactis* strain NZ9000 (14). Transformants were grown at 30°C in M17 medium

(Oxoid, Basingstoke, United Kingdom), supplemented with 1.0% (wt/vol) glucose, and 5 µg/ml chloramphenicol. Nisin was added when the optical density at 600 nm reached 0.7 (14), and cells were harvested 8 to 15 h later. Sodium dodecyl phosphate-polyacrylamide gel electrophoresis (SDS-PAGE) analysis of membrane fractions prepared by differential centrifugation (23) showed that FolT and ThiT were abundantly expressed (Fig. 2A) and had apparent molecular masses (18 and 22 kDa, respectively) near those predicted. Cells expressing FolT or ThiT, and empty-vector controls, were assayed for binding of ³H-labeled folates or thiamine after de-energization with 2-deoxyglucose to suppress interference by endogenous uptake systems (Fig. 2B to E). Cells expressing FolT bound large amounts of (6S)-[³H]folinic acid or [³H]folic acid (~17 pmol/mg protein), and those expressing ThiT bound a similar amount of [³H]thiamine. Adding a polyglutamyl tail of 2 to 4 residues to [³H]folic acid (16) markedly reduced binding, indicating that polyglutamyl folates are poor substrates for FolT, which is consistent with results from experiments using *L. casei* cells (22). In all cases, vitamin binding approached a plateau within 5 s and was rapidly reversed by adding an excess of unlabeled substrate. The observed vitamin acquisition, thus, has the characteristics of a binding process rather than those of an uptake process.

For further characterization, FolT and ThiT were tagged with N-terminal His₈ sequences. FolT-His and ThiT-His were produced in *L. lactis* as described above, except that cells were cultured in chemically defined medium (17, 19) without folic acid (for FolT-His) or thiamine (for ThiT-His) and harvested 3 h after induction. Membrane vesicles were prepared (24), and proteins were solubilized with dodecyl-β-D-maltoside (DDM) and purified to homogeneity by using nickel-Sepharose and gel filtration chromatography (3) (Fig. 3A and B). Vitamin binding was measured via quenching of intrinsic tryptophan fluorescence, using a Spex Fluorolog 322 spectrofluorometer (Jobin Yvon) and a 1-ml stirred cuvette at 25°C. The FolT-His and ThiT-His concentrations were 100 to 500 nM, and solutions of folinic acid, folic acid, or thiamine were added in 0.5- to 2-µl steps. Fluorescence was monitored at 340 nm for 20 to 30 s (excitation at 280 nm) after each substrate addition. Data were analyzed as described previously

* Corresponding author. Mailing address: University of Florida, Horticultural Sciences Department, P.O. Box 110690, Gainesville, FL 32611. Phone: (352) 392-1928. Fax: (352) 392-5653. E-mail: adha@ufl.edu.

† A.E. and G.B.E. contributed equally to the paper.

‡ Supplemental material for this article may be found at <http://j.b.asm.org/>.

∇ Published ahead of print on 5 September 2008.

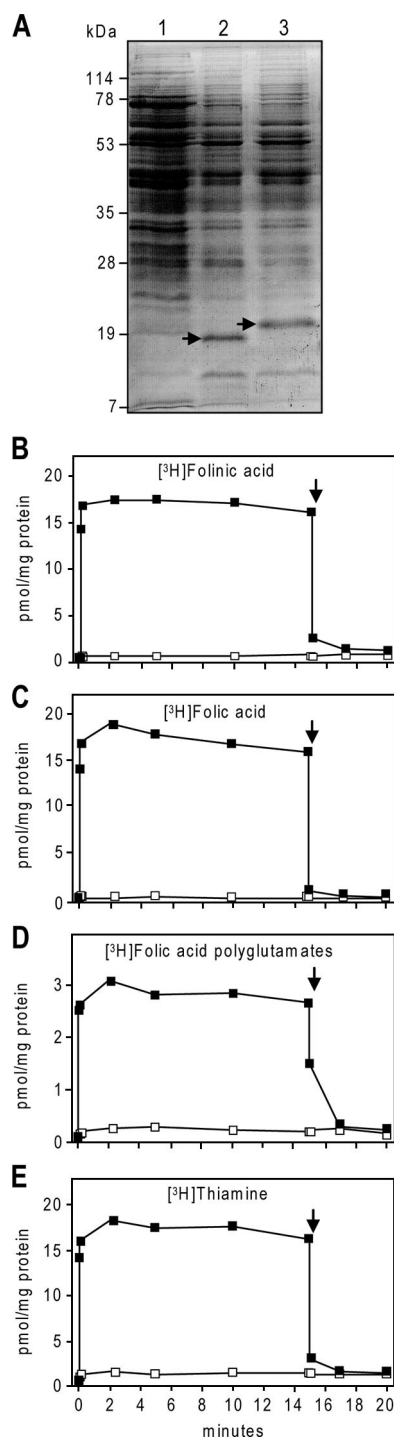


FIG. 2. Functional expression of *L. casei* FolT and ThiT in *L. lactis*. (A) SDS-PAGE (12% gel) of membrane fractions from *L. lactis* harboring pNZ8048 alone (lane 1; 50 μ g protein), or containing FolT (lane 2; 25 μ g protein) or ThiT (lane 3; 25 μ g protein). Staining was with Coomassie brilliant blue. The arrows indicate FolT and ThiT bands. Positions of molecular mass markers (kDa) are shown. (B to E) Binding of ^3H -labeled folates or thiamine to *L. lactis* cells harboring pNZ8048 alone (open squares) or expressing FolT or ThiT (filled squares). Assays (total volume, 1 ml) were performed in phosphate-buffered saline (PBS), pH 7.4, at 30°C with stirring. Cells were washed and resuspended (optical density at 600 nm, 20), and 0.5-ml aliquots were pretreated for 5 min with 2-deoxyglucose (25 mM final concentration). Assays were started by adding 0.5 ml of PBS containing

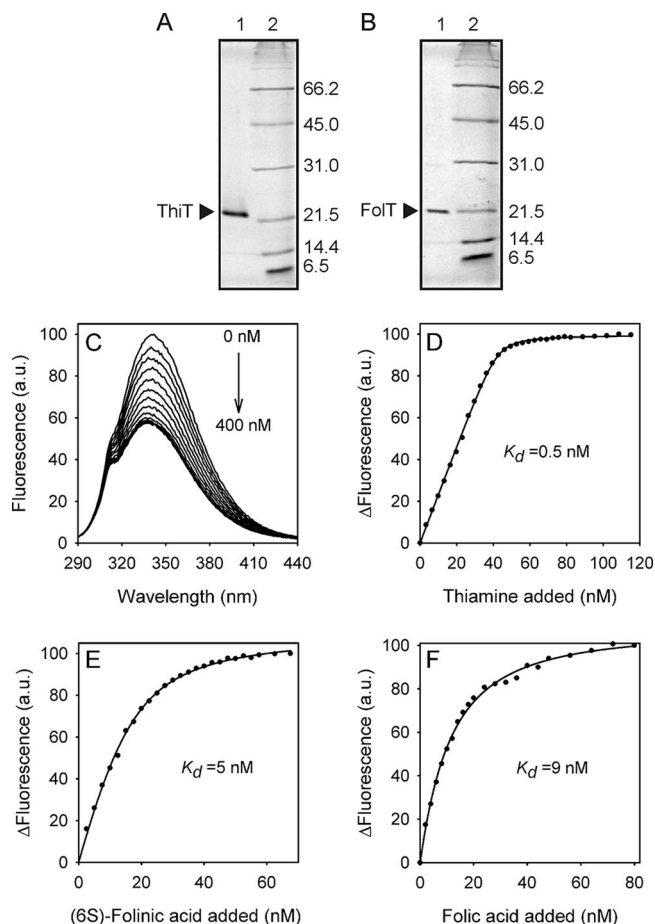


FIG. 3. Purification and characterization of His-tagged *L. casei* ThiT and FolT. (A and B) SDS-PAGE of purified ThiT-His and FolT-His, as in Fig. 2A. (C) Fluorescence spectrum of ThiT-His (320 nM in 50 mM K phosphate, 200 mM KCl, 0.05% [wt/vol] DDM, pH 7.0) in the absence of thiamine (uppermost trace) and in the presence of successively higher concentrations of thiamine (up to 400 nM). (D) Fluorescence titration of ThiT-His with thiamine. (E and F) Fluorescence titration of FolT-His (210 nM in 50 mM K phosphate, 200 mM KCl, 0.05% [wt/vol] DDM, pH 7.0) with (6S)-folic acid (E) and folic acid (F).

L. casei genome contains a gene cluster encoding homologs of BioN (LSEI_2472) and BioM (LSEI_2473 and LSEI_2474), which are thus candidates for shared components of the folate and thiamine transporters.

^3H -labeled vitamin (final concentration, 12.6 to 14.5 nM). At various times, cells (100 μ l) were harvested by vacuum filtration on a cellulose nitrate membrane (0.45 μ m). Filters were washed twice with 2 ml of ice-cold PBS, and their ^3H content was determined by scintillation counting. The arrows show when unlabeled vitamin was added to give a final concentration of 50 μ M. Cells expressing FolT were incubated with (6S)-[3',5',7,9- ^3H (N)]folic acid diammonium salt (Moravek; 10 Ci/mmol) (B), [3',5',7,9- ^3H]folic acid diammonium salt (Moravek; 25.9 Ci/mmol) (C), or [^3H]folic acid polyglutamates (45 Ci/mmol) comprising 40% tri-, 56% tetra-, and 4% pentaglutamates (D). Cells expressing ThiT were incubated with [^3H (G)]thiamine hydrochloride (ARC; 10 Ci/mmol) (E). ^3H -Labeled substrates were chromatographically purified before use (4, 13).

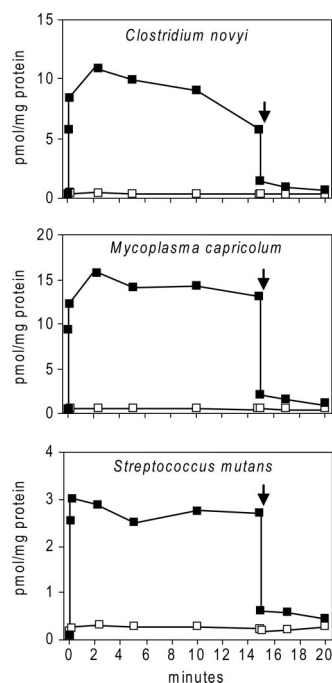


FIG. 4. Folate-binding by FoIT homologs from pathogenic Firmicutes expressed in *L. lactis*. The *folT* genes from *Clostridium novyi* and *Streptococcus mutans* were obtained by PCR from genomic DNA; that of *Mycoplasma capricolum* was synthesized by GenScript (Piscataway, NJ). Cells harboring pNZ8048 alone (open squares) or containing FoIT homologs (filled squares) were assayed for binding of (6S)-[³H]folinic acid (final concentration, 13.5 nM) as in Fig. 2. The arrows show when unlabeled folinic acid was added to give a final concentration of 50 μ M.

We thank Robert Burne (University of Florida) for *Streptococcus mutans* genomic DNA and Shibin Zhou (Johns Hopkins University School of Medicine) for *Clostridium novyi* genomic DNA.

This project was supported by National Institutes of Health grant R01 GM071382 (to A.D.H.), by The Netherlands Organization for Scientific Research (vidi grant to D.J.S.), and by an endowment from the C.V. Griffin, Sr. Foundation.

REFERENCES

- Davidson, A. L., E. Dassa, C. Orelle, and J. Chen. 2008. Structure, function, and evolution of bacterial ATP-binding cassette systems. *Microbiol. Mol. Biol. Rev.* **72**:317–364.
- de Crécy-Lagard, V., B. El Yacoubi, R. D. de la Garza, A. Noiriel, and A. D. Hanson. 2007. Comparative genomics of bacterial and plant folate synthesis and salvage: predictions and validations. *BMC Genomics* **8**:245.
- Duurkens, R. H., M. B. Tol, E. R. Geertsma, H. P. Permentier, and D. J. Slotboom. 2007. Flavin binding to the high affinity riboflavin transporter RibU. *J. Biol. Chem.* **282**:10380–10386.
- Gregory, J. F., III, and J. P. Toth. 1988. Chemical synthesis of deuterated folate monoglutamate and in vivo assessment of urinary excretion of deuterated folates in man. *Anal. Biochem.* **170**:94–104.
- Hebbeln, P., D. A. Rodionov, A. Alfandega, and T. Eitinger. 2007. Biotin uptake in prokaryotes by solute transporters with an optional ATP-binding cassette-containing module. *Proc. Natl. Acad. Sci. USA* **104**:2909–2914.
- Henderson, G. B., J. M. Kojima, and H. P. Kumar. 1985. Differential interaction of cations with the thiamine and biotin transport proteins of *Lactobacillus casei*. *Biochim. Biophys. Acta* **14**:201–206.
- Henderson, G. B., and S. Potuznik. 1982. Cation-dependent binding of substrate to the folate transport protein of *Lactobacillus casei*. *J. Bacteriol.* **150**:1098–1102.
- Henderson, G. B., and E. M. Zevely. 1978. Binding and transport of thiamine by *Lactobacillus casei*. *J. Bacteriol.* **133**:1190–1196.
- Henderson, G. B., E. M. Zevely, and F. M. Huennekens. 1977. Purification and properties of a membrane-associated, folate-binding protein from *Lactobacillus casei*. *J. Biol. Chem.* **252**:3760–3765.
- Henderson, G. B., E. M. Zevely, and F. M. Huennekens. 1979. Mechanism of folate transport in *Lactobacillus casei*: evidence for a component shared with the thiamine and biotin transport systems. *J. Bacteriol.* **137**:1308–1314.
- Henderson, G. B., E. M. Zevely, and F. M. Huennekens. 1979. Coupling of energy to folate transport in *Lactobacillus casei*. *J. Bacteriol.* **139**:552–559.
- Henderson, G. B., E. M. Zevely, R. J. Kadner, and F. M. Huennekens. 1977. The folate and thiamine transport proteins of *Lactobacillus casei*. *J. Supramol. Struct.* **6**:239–247.
- Kaushik, S., and K. S. Alexander. 2003. A modified reverse-phase HPLC method for the analysis of mexiletine hydrochloride. *J. Liq. Chromatogr.* **26**:1287–1296.
- Kuipers, O. P., P. G. de Ruyter, M. Kleerebezem, and W. M. de Vos. 1998. Quorum sensing-controlled gene expression in lactic acid bacteria. *J. Biotechnol.* **64**:15–21.
- Melnick, J., E. Lis, J.-H. Park, C. Kinsland, H. Mori, T. Baba, J. Perkins, G. Schyns, O. Vassieva, A. Osterman, and T. P. Begley. 2004. Identification of the two missing bacterial genes involved in thiamine salvage: thiamine pyrophosphokinase and thiamine kinase. *J. Bacteriol.* **186**:3660–3662.
- Naponelli, V., A. D. Hanson, and J. F. Gregory III. 2007. Improved methods for the preparation of [³H]folate polyglutamates: biosynthesis with *Lactobacillus casei* and enzymatic synthesis with *Escherichia coli* folylpolyglutamate synthetase. *Anal. Biochem.* **371**:127–134.
- Otto, R., B. H. ten Brink, H. Veldkamp, and W. N. Konings. 1983. The relation between growth rate and electrochemical proton gradient of *Streptococcus cremoris*. *FEMS Microbiol. Lett.* **16**:69–74.
- Overbeek, R., T. Begley, R. M. Butler, J. V. Choudhuri, H. Y. Chuang, M. Cohoon, V. de Crécy-Lagard, N. Diaz, T. Disz, R. Edwards, M. Fonstein, E. D. Frank, S. Gerdes, E. M. Glass, A. Goemann, A. Hanson, D. Iwata-Reuyl, R. Jensen, N. Jamshidi, L. Krause, M. Kubal, N. Larsen, B. Linke, A. C. McHardy, F. Meyer, H. Neweger, G. Olsen, R. Olson, A. Osterman, V. Portnoy, G. D. Pusch, D. A. Rodionov, C. Ruckert, J. Steiner, R. Stevens, I. Thiele, O. Vassieva, Y. Ye, O. Zagnitko, and V. Vonstein. 2005. The subsystems approach to genome annotation and its use in the project to annotate 1,000 genomes. *Nucleic Acids Res.* **33**:5691–5702.
- Poolman, B., and W. N. Konings. 1988. Relation of growth of *Streptococcus lactis* and *Streptococcus cremoris* to amino acid transport. *J. Bacteriol.* **170**:700–707.
- Rodionov, D. A., A. G. Vitreschak, A. A. Mironov, and M. S. Gelfand. 2002. Comparative genomics of thiamin biosynthesis in prokaryotes. New genes and regulatory mechanisms. *J. Biol. Chem.* **277**:48949–48959.
- Rodionov, D. A., P. Hebbeln, M. S. Gelfand, and T. Eitinger. 2006. Comparative and functional genomic analysis of prokaryotic nickel and cobalt uptake transporters: evidence for a novel group of ATP-binding cassette transporters. *J. Bacteriol.* **188**:317–327.
- Shane, B., and E. L. Stokstad. 1975. Transport and metabolism of folates by bacteria. *J. Biol. Chem.* **250**:2243–2253.
- Valyasevi, R., W. E. Sandine, and B. Geller. 1991. A membrane protein is required for bacteriophage ϕ 2 infection of *Lactococcus lactis* subsp. *lactis* C2. *J. Bacteriol.* **173**:6095–6100.
- van der Heide, T., and B. Poolman. 2000. Osmoregulated ABC-transport system of *Lactococcus lactis* senses water stress via changes in the physical state of the membrane. *Proc. Natl. Acad. Sci. USA* **97**:7102–7106.
- Veldhuis, G., E. P. Vos, J. Broos, B. Poolman, and R. M. Scheek. 2004. Evaluation of the flow-dialysis technique for analysis of protein-ligand interactions: an experimental and a Monte Carlo study. *Biophys. J.* **86**:1959–1968.
- Wang, P., N. Nirmalan, Q. Wang, P. F. Sims, and J. E. Hyde. 2004. Genetic and metabolic analysis of folate salvage in the human malaria parasite *Plasmodium falciparum*. *Mol. Biochem. Parasitol.* **135**:77–87.
- Wilkins, M. R., C. Pasquali, R. D. Appel, K. Ou, O. Golaz, J. C. Sanchez, J. X. Yan, A. A. Gooley, G. Hughes, I. Humphery-Smith, K. L. Williams, and D. F. Hochstrasser. 1996. From proteins to proteomes: large scale protein identification by two-dimensional electrophoresis and amino acid analysis. *Bio/Technology* **14**:61–65.
- Winkler, W., A. Nahvi, and R. R. Breaker. 2002. Thiamine derivatives bind messenger RNAs directly to regulate bacterial gene expression. *Nature* **419**:952–956.