A *Legionella pneumophila* Peptidyl-Prolyl *cis*-*trans* Isomerase Present in Culture Supernatants Is Necessary for Optimal Growth at Low Temperatures

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Several *Legionella pneumophila* **proteins were highly expressed in low-temperature supernatants. One of these proteins was the peptidyl-prolyl isomerase PpiB. Mutants lacking** *ppiB* **exhibited reduced growth at 17°C. Since PpiB lacked a signal sequence and was present in 17°C supernatants of type II and type IV secretion mutants, this protein may be secreted by a novel mechanism.**

Legionella pneumophila is the agent of Legionnaires' disease pneumonia (14), and infection results from inhalation of contaminated water droplets from aerosol-generating devices (28). *L. pneumophila* occurs naturally in freshwaters (3, 15, 23, 29, 40), but it is also widespread in man-made water systems (26, 33, 45). It exists in the planktonic phase, in biofilms, and as an intracellular parasite of protozoans (12, 24, 25, 29, 31). But the distribution of *L. pneumophila* is also likely due to its capacity to survive at 4 to 63° C (15, 23, 44, 45). Thus, there is need for understanding *L. pneumophila* survival at low temperatures. Recently, we observed that *L. pneumophila lsp* mutants deficient in type II secretion grow normally at 30 to 37°C but their growth at 12 to 25°C is impaired (6, 22, 41). The wild type stimulates the growth of an *lsp* mutant at 25°C when they are plated near each other (41), suggesting that secreted factors promote low-temperature growth.

To identify secreted proteins that are newly expressed or hyperexpressed when the wild type is grown at low temperatures, strain $130b$ (= ATCC BAA-74) was grown to late log phase in buffered yeast extract (BYE) broth at 37, 17, and 12°C, and then filter-sterilized culture supernatants were examined by two-dimensional polyacrylamide gel electrophoresis, as previously described (10). Although growth slowed with decreasing temperature, there were many similarities between the profiles (Fig. 1), indicating that *L. pneumophila* secretes proteins while it is growing at 12 to 37°C. Low-temperature supernatants had slightly fewer proteins rather than more proteins, indicating that low-temperature incubation does not result in wholesale lysis. In addition to the similarities, there were differences. For example, the amounts of some proteins present at 37°C were reduced at 17 to 12°C (Fig. 1). There were also proteins that were more pronounced in the 17 and 12°C samples (e.g., spots 1 to 4 in Fig. 1). Thus, *L. pneumophila* secretion changes when the organisms is grown at low temperatures. We hypothesized that proteins whose amounts are

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greater at 17 to 12°C contribute to survival at low temperatures.

To identify such proteins, spots 1 and 2 were analyzed by mass spectrometry (10), and the data were compared to a database (http://genolist.pasteur.fr/LegioList/). The 164-amino-acid protein in spot 1 was identified as PpiB (Lcy), a peptidyl-prolyl isomerase (PPIase) belonging to the cyclophilin family. *ppiB* is either monocistronic or the last gene in a small operon, and the preceding open reading frame encodes a tRNA ribosyltransferase. The 188-amino-acid protein in spot 2 was also annotated as a cyclophilin PPIase, and in the sequenced strains Philadelphia, Paris, and Lens the monocistronic genes that encode it are lpg1962, lpp1946, and lpl1936, respectively. Previously, PpiB was purified from cytoplasmic extracts of *L. pneumophila* Philadelphia-1 and demonstrated to have PPIase activity (37). In that study, however, there was no attempt to look for protein in supernatants or in bacteria grown at temperatures other than 37°C. There have been no previous studies of the putative PPIase of spot 2; thus, we refer to this protein as Lpg1962. Because PpiB and Lpg1962 were in wild-type supernatants that did not show evidence of cell lysis or leakage, these two proteins likely are secreted proteins. The fact that a previous study had shown that PpiB was present in cell extracts does not invalidate our findings. For example, cell-associated PpiB might simply represent a protein on its way toward secretion, or perhaps PpiB is maintained within and outside the cell. Alternatively, this protein might exist mainly in cells at 37°C but be secreted at 17°C. Lpg1962 has a signal sequence (16, 27) and thus is likely a substrate for type II secretion. PpiB lacks a typical signal sequence or a signal sequence with twin arginines (34), suggesting that its presence in supernatants depends on another mechanism.

To determine if PpiB and Lpg1962 promote growth at low temperatures, we used allelic exchange to construct mutants. First, primers were designed to amplify genes from 130b DNA; Mas23 (5'-CGTACGGAGCTCATATTCAG) and Mas25 (5'-TGGTAATATTTTCAATGACTACAGG) yielded a 773-bp fragment for lpg1962, and Mas26 (5-TCTGCAATGAATAC GGATGG) and Mas27 (5-GGTACA CAAAAAGTTCT CGC) yielded a 1,552-bp fragment containing *ppiB*. Fragments containing *ppiB* and lpg1962 were ligated into pGEM-T Easy, yielding pB24 and pR11. pB24 was digested with BamHI,

FIG. 1. Two-dimensional polyacrylamide gel electrophoresis analysis of culture supernatants obtained from *L. pneumophila* grown at low temperatures. Wild-type strain 130b was grown in BYE broth to late log phase at 37, 17, or 12°C, and then the proteins in the different cell-free culture supernatants were separated by two-dimensional polyacrylamide gel electrophoresis and stained with Coomassie blue. Some protein spots whose amounts were greater in the 37°C sample are indicated by letters in the left panel, whereas some of the proteins that were more prominently expressed in the low-temperature samples are indicated by numbers in the center and right panels. Similar protein profiles were obtained on at least two other occasions using material derived from independent cultures.

which cut 271 bp after the *ppiB* start codon, and was then ligated to a Kmr gene from pMB2190 (17) to obtain pB24K or to a Gm^r gene from pX1918GT (39) to produce pB24G2. Next, NotI fragments of pB24K and pB24G2 containing the disrupted genes were cloned into the SmaI site of pRE112 (13) to obtain pB24KS3 and pB24GS4. pR11 was digested with AgeI, which cut 292 bp after the lpg1962 start, and then was ligated to the Km^r and Gm^r cassettes, resulting in pR11K1 and pR11G2. Following NotI digestion, the disrupted lpg1962 genes were cloned into pRE112, yielding Km^r pR11K1S3 and Gm^r pR11G2S3. 130b was transformed with pR11G2S3, pR11K1S3, pB24GS4, and pB24KS3 by electroporation (7), and mutants were selected as previously described (34). To construct lpg1962 *ppiB* double mutants, a Gm^r lpg1962 mutant was transformed with pB24KS3, and a Km^r lpg1962 mutant was transformed with pB24GS4. Ultimately, six mutants were obtained: Gmr NU340 and Kmr NU341 for *ppiB*, Gmr NU342 and Kmr NU343 for lpg1962, and Kmr Gmr NU344 and NU345 for *ppiB* lpg1962. All mutants grew normally on buffered charcoal-yeast extract (BCYE) agar and in BYE broth at 37°C (Fig. 2A; data not shown), indicating that *ppiB* and lpg1962 are not required for extracellular growth under standard conditions. The fact that a *ppiB* mutant grows normally at 37°C was previously observed (37).

Next, we compared the growth of 130b and the growth of the mutants on BCYE agar at 17°C (Fig. 2A). As hypothesized, *ppiB* mutant NU340 displayed reduced survival at this low temperature. Since independent *ppiB* mutant NU341 also showed impaired growth (data not shown), these data indicated that the defect was due to the loss of *ppiB* rather than second-site mutations. When an intact *ppiB* gene was reintroduced into NU340 on pMB3, the wild type and the *ppiB* mutant grew comparably (Fig. 2B). To create pMB3, *ppiB* was amplified using Mas27 and Mas28 (5'-TGTTTTGCATGATG TTTGTAAT) and cloned into pMMB2002 (35). In contrast to the results for *ppiB*, lpg1962 mutants NU342 and NU343 grew like the wild type when they were plated at low temperature (Fig. 2A; data not shown). Compatible with these results, the *ppiB* lpg1962 double mutants had reduced abilities to grow in the same way as the *ppiB* mutants (Fig. 2A; data not shown). To confirm the results obtained by plating, we compared the growth of 130b and the growth of mutants in BYE broth at 17°C. The *ppiB* mutants and the double mutants exhibited reduced growth, whereas lpg1962 mutants grew normally (Fig. 3A). After reintroduction of *ppiB*, the wild type and the *ppiB* mutants grew comparably in broth at 17°C (Fig. 3B). These data indicate that PpiB, but not Lpg1962, is necessary for optimal extracellular growth at low temperatures. The *ppiB* mutants were not as impaired at low temperatures as *lspF* mutant NU275 (Fig. 2A and 3A) (35, 41), suggesting that additional secreted factors promote low-temperature growth. Indeed, a *ppiB* mutant was able to stimulate the growth of the *lspF* mutant at a low temperature when these mutants were plated near each other (data not shown).

Because of the newfound importance of PpiB, we further investigated the mechanism by which this protein appears in supernatants. As noted previously, the absence of a Sec- or Tat-dependent signal sequence indicated that there is not type II secretion. Compatible with this, supernatants from an *lspF* mutant grown at 17°C contained PpiB (Fig. 4). But the presence of more proteins in the mutant supernatants than in wild-type supernatants (Fig. 1), rather than many fewer proteins, as observed when 37°C supernatants were compared (10), suggests that a type II mutant undergoes a greater degree of leakage or lysis at 17°C. *L. pneumophila* also possesses Lvh type IVA and Dot/Icm type IVB secretion (42). Proteins exported via type IV secretion often do not contain typical signal sequences (18). Thus, we examined supernatants obtained from *dotG* mutant AA405 and *lvhB9* mutant AA474 (from Cary Engleberg, University of Michigan) grown at 17°C (Fig. 4). In both cases, PpiB was present, indicating that the PPIase does not require one of the type IV pathways for export. Neither type IV mutant exhibited reduced growth on BCYE agar or in BYE broth at 17°C (data not shown). These data suggest that PpiB is not released by one of the three known *Legionella* secretion systems. Genome sequencing has suggested the presence of a type I secretion system in *L. pneumophila*, as well as the presence of type V secretion in some strains (4, 21). However, PpiB lacks the glycine-rich repeats often present in type I substrates (11), and the "autotransport-

FIG. 2. Low-temperature growth of the wild type and *ppiB* and lpg1962 mutants of *L. pneumophila* on BCYE agar. (A) Tenfold serial dilutions of wild-type strain 130b, *ppiB* mutant NU340, lpg1962 mutant NU342, *ppiB* lpg1962 double mutant NU344, and *lspF* mutant NU275 containing equivalent numbers of CFU (as determined at 37°C) were spotted onto BCYE agar plates and then incubated at either 37°C (left panel) or 17°C (right panel). Images of bacterial growth were then obtained after 3 days for the 37°C plates and after 20 days for the 17°C plates. The results are representative of at least three independent experiments. (B) Tenfold serial dilutions of wild-type strain 130b and *ppiB* mutant NU340 containing either the vector pMMB2002 or the complementing plasmid pMB3 (i.e., *ppiB* cloned into pMMB2002) were spotted onto BCYE agar and then incubated at 37°C (left panel) or 17°C (right panel). Images of bacterial growth were then obtained as described above. For unknown reasons, the vector alone slowed the growth of both the wild type and the mutant at 17°C (compared to the 17°C cultures used for panel A). Nonetheless, the specific growthstimulating effect of cloned *ppiB* on NU340 at 17°C is still evident. The results are representative of two independent experiments.

ers" of type V secretion generally contain Sec-dependent signal sequences (18). *L. pneumophila* lacks type III secretion but does express flagella, which might provide a pathway for export (8, 20).

Regardless of variations in the localization of PpiB or its secretion mechanism, our analyses indicate that PpiB is required for optimal growth at low temperatures. Given its PPIase activity (37), PpiB may catalyze the isomerization of secreted proteins, such as type II substrates, that promote survival at low temperatures. Alternatively, PpiB might assist cold-adapted exoenzymes as a chaperone (5, 19, 36, 38). Finally, PpiB, whether as an isomerase or as a chaperone, might promote the functioning of a secretion apparatus. There are examples of PPIases that are secreted and surface expressed by microbes, such as

FIG. 3. Low-temperature growth of the wild type and *ppiB* and lpg1962 mutants of *L. pneumophila* in BYE broth. Log-phase bacteria were inoculated into BYE broth at 17°C, and then the growth of the indicated strains was monitored by recording the optical densities of the cultures at various times. (A) Comparison of the growth of wildtype strain 130b (\bullet), *ppiB* mutant NU340 (\circ), lpg1962 mutant NU342 $(\vec{\nabla})$, *ppiB* lpg1962 double mutant NU344 ($\vec{\nabla}$), and *lspF* mutant NU275 (\blacksquare) . (B) Comparison of the growth of wild-type strain 130b (\blacksquare) , *ppiB* mutant NU340 (\circ), and complemented *ppiB* mutant NU340 ($pMB3$) (\Box) . The apparent differences in growth between either the *ppiB* mutant or the *ppiB* lpg1962 mutant and the wild-type strain or the complemented *ppiB* mutant were statistically significant, as were the differences in growth between the *lspF* mutant and the other strains $(P < 0.05$, Student's *t* test). The slight differences in optical densities at 660 nm (OD660) between the wild type and the lpg1962 mutant observed at some time points were not seen in repeat experiments. The data are the means and standard deviations for duplicate cultures and are representative of at least three (A) and two (B) independent experiments.

HP0175 of *Helicobacter pylori* and Mip of *L. pneumophila* (1, 9, 30), and there are PPIases that have ben previously linked to low-temperature adaptation, such as cell-associated PpiB of *Bacillus subtilis*, FKBP of *Shewanella* sp., and RotA of *Erwinia chrysanthemi* (2, 32, 43). But the connection that we uncovered between a secreted PPIase and low-temperature growth is a novel observation.

Previously, Schmidt et al. observed that a *ppiB* mutant has a reduced ability to grow in *Acanthamoeba castellanii*, an aquatic protozoan that serves as an intracellular niche for *L. pneumophila* (37). Thus, our data for the extracellular growth of *ppiB* mutants at low temperature indicate that PpiB likely promotes survival in natural habitats by at least two mechanisms. For

FIG. 4. Two-dimensional polyacrylamide gel electrophoresis analysis of culture supernatants obtained from *L. pneumophila* type II and type IV secretion mutants grown at low temperatures. *lspF* mutant NU275, *dotG* mutant AA405, and *lvhB9* mutant AA474 were grown in BYE broth to late log phase at 17°C, and then the proteins in the cell-free culture supernatants were separated by two-dimensional polyacrylamide gel electrophoresis and stained with Coomassie blue. The PpiB spot is indicated by an arrow.

Lpg1962, the absence of a growth defect in the lpg1962 mutant does not necessarily indicate irrelevance for low-temperature growth, since it is possible that another PPIase can replace Lpg1962. Combined, the present findings involving PpiB and previous work on Lsp indicate that a variety of secretion functions aid *L. pneumophila*, and perhaps other bacteria, in growing at low temperatures.

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REFERENCES

- 1. **Basak, C., S. K. Pathak, A. Bhattacharyya, S. Pathak, J. Basu, and M. Kundu.** 2005. The secreted peptidyl prolyl *cis*,*trans*-isomerase HP0175 of *Helicobacter pylori* induces apoptosis of gastric epithelial cells in a TLR4- and apoptosis signal-regulating kinase 1-dependent manner. J. Immunol. **174:** 5672–5680.
- 2. **Budde, I., L. Steil, C. Scharf, U. Volker, and E. Bremer.** 2006. Adaptation of *Bacillus subtilis* to growth at low temperature: a combined transcriptomic and proteomic appraisal. Microbiology **152:**831–853.
- 3. **Carvalho, F. R., R. F. Vazoller, A. S. Foronda, and V. H. Pellizari.** 2007. Phylogenetic study of *Legionella* species in pristine and polluted aquatic samples from a tropical Atlantic forest ecosystem. Curr. Microbiol. **55:**288– 293.
- 4. **Cazalet, C., C. Rusniok, H. Bruggemann, N. Zidane, A. Magnier, L. Ma, M. Tichit, S. Jarraud, C. Bouchier, F. Vandenesch, F. Kunst, J. Etienne, P. Glaser, and C. Buchrieser.** 2004. Evidence in the *Legionella pneumophila* genome for exploitation of host cell functions and high genome plasticity. Nat. Genet. **36:**1165–1173.
- 5. **Chakraborty, A., B. Sen, R. Datta, and A. K. Datta.** 2004. Isomerase-independent chaperone function of cyclophilin ensures aggregation prevention of adenosine kinase both *in vitro* and under *in vivo* conditions. Biochemistry **43:**11862–11872.
- 6. **Cianciotto, N. P.** 2005. Type II secretion: a protein secretion system for all seasons. Trends Microbiol. **13:**581–588.
- 7. **Cianciotto, N. P., and B. S. Fields.** 1992. *Legionella pneumophila mip* gene potentiates intracellular infection of protozoa and human macrophages. Proc. Natl. Acad. Sci. USA **89:**5188–5191.
- 8. **Cornelis, G. R.** 2006. The type III secretion injectisome. Nat. Rev. Microbiol. **4:**811–825.
- 9. **DebRoy, S., V. Aragon, S. Kurtz, and N. P. Cianciotto.** 2006. *Legionella pneumophila* Mip, a surface-exposed peptidylproline *cis-trans*-isomerase, promotes the presence of phospholipase C-like activity in culture supernatants. Infect. Immun. **74:**5152–5160.
- 10. **DebRoy, S., J. Dao, M. Soderberg, O. Rossier, and N. P. Cianciotto.** 2006. *Legionella pneumophila* type II secretome reveals unique exoproteins and a chitinase that promotes bacterial persistence in the lung. Proc. Natl. Acad. Sci. USA **103:**19146–19151.
- 11. **Delepelaire, P.** 2004. Type I secretion in gram-negative bacteria. Biochim. Biophys. Acta **1694:**149–161.
- 12. **Donlan, R. M., T. Forster, R. Murga, E. Brown, C. Lucas, J. Carpenter, and B. Fields.** 2005. *Legionella pneumophila* associated with the protozoan *Hart-*

mannella vermiformis in a model multi-species biofilm has reduced susceptibility to disinfectants. Biofouling **21:**1–7.

- 13. **Edwards, R. A., L. H. Keller, and D. M. Schifferli.** 1998. Improved allelic exchange vectors and their use to analyze 987P fimbria gene expression. Gene **207:**149–157.
- 14. **Fields, B. S., R. F. Benson, and R. E. Besser.** 2002. *Legionella* and Legionnaires' disease: 25 years of investigation. Clin. Microbiol. Rev. **15:**506–526.
- 15. **Fliermans, C. B., W. B. Cherry, L. H. Orrison, S. J. Smith, D. L. Tison, and D. H. Pope.** 1981. Ecological distribution of *Legionella pneumophila*. Appl. Environ. Microbiol. **41:**9–16.
- 16. **Gardy, J. L., M. R. Laird, F. Chen, S. Rey, C. J. Walsh, M. Ester, and F. S. Brinkman.** 2005. PSORTb v. 2.0: expanded prediction of bacterial protein subcellular localization and insights gained from comparative proteome analysis. Bioinformatics **21:**617–623.
- 17. **Grindley, N. D., and C. M. Joyce.** 1980. Genetic and DNA sequence analysis of the kanamycin resistance transposon Tn*903*. Proc. Natl. Acad. Sci. USA **77:**7176–7180.
- 18. **Henderson, I. R., F. Navarro-Garcia, M. Desvaux, R. C. Fernandez, and D. Ala'Aldeen.** 2004. Type V protein secretion pathway: the autotransporter story. Microbiol. Mol. Biol. Rev. **68:**692–744.
- 19. **Hennecke, G., J. Nolte, R. Volkmer-Engert, J. Schneider-Mergener, and S. Behrens.** 2005. The periplasmic chaperone SurA exploits two features characteristic of integral outer membrane proteins for selective substrate recognition. J. Biol. Chem. **280:**23540–23548.
- 20. **Heuner, K., and M. Steinert.** 2003. The flagellum of *Legionella pneumophila* and its link to the expression of the virulent phenotype. Int. J. Med. Microbiol. **293:**133–143.
- 21. **Jacobi, S., and K. Heuner.** 2003. Description of a putative type I secretion system in *Legionella pneumophila*. Int. J. Med. Microbiol. **293:**349–358.
- 22. **Johnson, T. L., J. Abendroth, W. G. Hol, and M. Sandkvist.** 2006. Type II secretion: from structure to function. FEMS Microbiol. Lett. **255:**175–186.
- 23. **Joly, J. R., M. Boissinot, J. Duchaine, M. Duval, J. Rafrafi, D. Ramsay, and R. Letarte.** 1984. Ecological distribution of *Legionellaceae* in the Quebec City area. Can. J. Microbiol. **30:**63–67.
- 24. **Mampel, J., T. Spirig, S. S. Weber, J. A. Haagensen, S. Molin, and H. Hilbi.** 2006. Planktonic replication is essential for biofilm formation by *Legionella pneumophila* in a complex medium under static and dynamic flow conditions. Appl. Environ. Microbiol. **72:**2885–2895.
- 25. **Marrao, G., A. Verissimo, R. G. Bowker, and M. S. daCosta.** 1993. Biofilms as major sources of *Legionella* spp. in hydrothermal areas and their dispersion into stream water. FEMS Microbiol. Ecol. **12:**25–33.
- 26. **Mouchtouri, V., E. Velonakis, A. Tsakalof, C. Kapoula, G. Goutziana, A. Vatopoulos, J. Kremastinou, and C. Hadjichristodoulou.** 2007. Risk factors for contamination of hotel water distribution systems by *Legionella* species. Appl. Environ. Microbiol. **73:**1489–1492.
- 27. **Nielsen, H., J. Engelbrecht, S. Brunak, and G. von Heijne.** 1997. Identification of prokaryotic and eukaryotic signal peptides and prediction of their cleavage sites. Protein Eng. **10:**1–6.
- 28. **O'Loughlin, R. E., L. Kightlinger, M. C. Werpy, E. Brown, V. Stevens, C. Hepper, T. Keane, R. F. Benson, B. S. Fields, and M. R. Moore.** 2007. Restaurant outbreak of Legionnaires' disease associated with a decorative fountain: an environmental and case-control study. BMC Infect. Dis. **7:**93– 101.
- 29. **Paszko-Kolva, C., M. Shahamat, and R. R. Colwell.** 1993. Effect of temperature on survival of *Legionella pneumophila* in the aquatic environment. Microb. Releases **2:**73–79.
- 30. **Pereira, P. J., M. C. Vega, E. Gonzalez-Rey, R. Fernandez-Carazo, S. Macedo-Ribeiro, F. X. Gomis-Ruth, A. Gonzalez, and M. Coll.** 2002.

Trypanosoma cruzi macrophage infectivity potentiator has a rotamase core and a highly exposed alpha-helix. EMBO Rep. **3:**88–94.

- 31. **Piao, Z., C. C. Sze, O. Barysheva, K. Iida, and S. Yoshida.** 2006. Temperature-regulated formation of mycelial mat-like biofilms by *Legionella pneumophila*. Appl. Environ. Microbiol. **72:**1613–1622.
- 32. **Pissavin, C., and N. Hugouvieux-Cotte-Pattat.** 1997. Characterization of a periplasmic peptidyl-prolyl *cis-trans* isomerase in *Erwinia chrysanthemi*. FEMS Microbiol. Lett. **157:**59–65.
- 33. **Ragull, S., M. Garcia-Nunez, M. L. Pedro-Botet, N. Sopena, M. Esteve, R. Montenegro, and M. Sabria.** 2007. *Legionella pneumophila* in cooling towers: fluctuations in counts, determination of genetic variability by pulsed-field gel electrophoresis (PFGE), and persistence of PFGE patterns. Appl. Environ. Microbiol. **73:**5382–5384.
- 34. **Rossier, O., and N. P. Cianciotto.** 2005. The *Legionella pneumophila tatB* gene facilitates secretion of phospholipase C, growth under iron-limiting conditions, and intracellular infection. Infect. Immun. **73:**2020–2032.
- 35. **Rossier, O., S. Starkenburg, and N. P. Cianciotto.** 2004. *Legionella pneumophila* type II protein secretion promotes virulence in the A/J mouse model of Legionnaires' disease pneumonia. Infect. Immun. **72:**310–321.
- 36. **Saul, F. A., J. P. Arie, B. Vulliez-le Normand, R. Kahn, J. M. Betton, and G. A. Bentley.** 2004. Structural and functional studies of FkpA from *Escherichia coli*, a *cis/trans* peptidyl-prolyl isomerase with chaperone activity. J. Mol. Biol. **335:**595–608.
- 37. **Schmidt, B., T. Tradler, J. U. Rahfeld, B. Ludwig, B. Jain, K. Mann, K. P. Rucknagel, B. Janowski, A. Schierhorn, G. Kullertz, J. Hacker, and G. Fischer.** 1996. A cyclophilin-like peptidyl-prolyl cis/trans isomerase from *Legionella pneumophila*—characterization, molecular cloning and overexpression. Mol. Microbiol. **21:**1147–1160.
- 38. **Scholz, C., B. Eckert, F. Hagn, P. Schaarschmidt, J. Balbach, and F. X. Schmid.** 2006. SlyD proteins from different species exhibit high prolyl isomerase and chaperone activities. Biochemistry **45:**20–33.
- 39. **Schweizer, H. P., and T. T. Hoang.** 1995. An improved system for gene replacement and *xylE* fusion analysis in *Pseudomonas aeruginosa*. Gene **158:** 15–22.
- 40. **Sheehan, K. B., J. M. Henson, and M. J. Ferris.** 2005. *Legionella* species diversity in an acidic biofilm community in Yellowstone National Park. Appl. Environ. Microbiol. **71:**507–511.
- 41. Söderberg, M. A., O. Rossier, and N. P. Cianciotto. 2004. The type II protein secretion system of *Legionella pneumophila* promotes growth at low temperatures. J. Bacteriol. **186:**3712–3720.
- 42. **Steinert, M., K. Heuner, C. Buchrieser, C. Albert-Weissenberger, and G. Glockner.** 2007. *Legionella* pathogenicity: genome structure, regulatory networks and the host cell response. Int. J. Med. Microbiol. **297:**577–587.
- 43. **Suzuki, Y., M. Haruki, K. Takano, M. Morikawa, and S. Kanaya.** 2004. Possible involvement of an FKBP family member protein from a psychrotrophic bacterium *Shewanella* sp. SIB1 in cold-adaptation. Eur. J. Biochem. **271:**1372–1381.
- 44. **Wadowsky, R. M., R. Wolford, A. M. McNamara, and R. B. Yee.** 1985. Effect of temperature, pH, and oxygen level on the multiplication of naturally occurring *Legionella pneumophila* in potable water. Appl. Environ. Microbiol. **49:**1197–1205.
- 45. **Wullings, B. A., and D. van der Kooij.** 2006. Occurrence and genetic diversity of uncultured *Legionella* spp. in drinking water treated at temperatures below 15°C. Appl. Environ. Microbiol. **72:**157–166.