Expression of the Plague Plasminogen Activator in *Yersinia pseudotuberculosis* and *Escherichia coli*

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Enteropathogenic yersiniae (*Yersinia pseudotuberculosis* **and** *Yersinia enterocolitica***) typically cause chronic disease as opposed to the closely related** *Yersinia pestis***, the causative agent of bubonic plague. It is established that this difference reflects, in part, carriage by** *Y. pestis* **of a unique 9.6-kb pesticin or Pst plasmid (pPCP) encoding plasminogen activator (Pla) rather than distinctions between shared** ;**70-kb low-calcium-response, or Lcr, plasmids (pCD in** *Y. pestis* **and pYV in enteropathogenic yersiniae) encoding cytotoxic Yops and anti-inflammatory V antigen. Pla is known to exist as a combination of 32.6-kDa (**a**-Pla) and slightly smaller (**b**-Pla) outer membrane proteins, of which at least one promotes bacterial dissemination in vivo and degradation of Yops in vitro. We show here that only** a**-Pla accumulates in** *Escherichia coli* **LE392/pPCP1 cultivated in enriched medium and that either autolysis or extraction of this isolate with 1.0 M NaCl results in release of soluble** α and β forms possessing biological activity. This process also converted cell-bound α -Pla to β -Pla **and smaller forms in** *Y. pestis* **KIM/pPCP1 and** *Y. pseudotuberculosis* **PB1/**1**/pPCP1 but did not promote solubilization. Pla-mediated posttranslational hydrolysis of pulse-labeled Yops in** *Y. pseudotuberculosis* **PB1/** 1**/pPCP1 occurred more slowly than that in** *Y. pestis* **but was otherwise similar except for accumulation of stable degradation products of YadA, a pYV-mediated fibrillar adhesin not encoded in frame by pCD. Carriage of pPCP by** *Y. pseudotuberculosis* **did not significantly influence virulence in mice.**

Bubonic plague caused by *Yersinia pestis* is among the most severe acute bacterial diseases of man. In contrast, closely related *Yersinia pseudotuberculosis* and the more distant *Yersinia enterocolitica* (44) typically cause chronic enteropathogenic infections in humans. Both syndromes depend upon the ability to express a low-calcium response (LCR) at 37°C but not 26°C, mediated by \sim 70-kb shared Lcr plasmids (termed pCD in *Y. pestis* and pYV in enteropathogenic yersiniae). The LCR is characterized by either vegetative growth in vitro in the presence of \geq 2.5 mM Ca²⁺ accompanied by downregulation of LcrF-regulated virulence factors (VirF in *Y. enterocolitica*) or else bacteriostasis associated with upregulation of these same determinants during cultivation in Ca^{2+} -deficient (≤ 1.0) mM) medium (19, 47) (Lcr^+). Lcr plasmids promoting this phenotype are functionally interchangeable (67) and essentially identical (49, 50) except that that from *Y. pestis* possesses frameshifts preventing expression of the fibrillar outer membrane adhesin YadA (31) and the lipoprotein YlpA (17) encoded by pYV (26, 54). Major effectors of virulence encoded by both pCD and pYV include anti-inflammatory V antigen and the secreted cytotoxins YopE, YopH, and YopO (YpkA) (19, 47).

The unique ability of *Y. pestis* to promote acute disease is caused in part by accessory virulence factors mediated by two plasmids not shared by the enteropathogenic yersiniae (4, 22, 51). One of these is a 9.6 kb-pesticin or Pst plasmid (pPCP) which encodes the 39.9-kDa bacteriocin pesticin (3, 10, 27, 48, 52), its 16-kDa immunity protein (48, 52), and a particulate plasminogen activator (2) termed Pla $(Pst⁺)$. Pla facilitates dissemination of yersiniae in host tissues; thus, its presence is essential for expression of virulence via transmission by fleabite and other peripheral routes of infection (8). Invasion of tissue is accomplished by Pla-dependent adherence to host basement membrane and extracellular matrix, where plasminogen activation facilitates bacterial metastasis (35). A transient 34.6-kDa primary *pla* product is processed upon insertion into the outer membrane to yield 32.6 -kDa α -Pla (58), which can then be hydrolyzed in \sim 2 h (43) to yield a slightly smaller derivative, termed β -Pla (58); both forms, especially β -Pla, accumulate as major outer membrane proteins (43, 62). The apparent molecular mass of α -Pla (\sim 35 kDa) estimated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) exceeds the correct value determined by sequencing (59) ; this anomaly probably also occurs for β -Pla $\overline{(-33 \text{ kDa by SDS-PAGE)}}$.

Native Yops did not accumulate in wild-type cells of *Y. pestis* grown in a \bar{Ca}^{2+} -deficient environment that favored their net synthesis by *Y. pseudotuberculosis* (61), although analysis of unprocessed Lcr⁺ cells of *Y. pestis* grown in vitro revealed a large segment of YopE (13) . Upon pulsing Ca²⁺-starved plague bacilli with $[355]$ methionine, radioactivity immediately appeared in a complete complement of native Yops but was promptly chased into small peptides (43, 56), except for a stable fragment of YopE probably identical to that noted above (13). This posttranslational degradation was attributed to the action of Pla (56, 57), a conclusion later verified by genetic analysis (60). These findings illustrate that Yops are cryptic in Lcr⁺ Pst⁺ isolates of *Y. pestis* grown in vitro but that they are obviously synthesized and delivered in an intact form in vivo because their structural genes are essential for the expression of disease (19, 47).

Essentially nothing is known about the mechanism whereby Pla mediates hydrolysis of Yops synthesized by *Y. pestis* or the

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Cloning vector

 $pPCP::Tn1$ (insert in *pst*)

Plasmids

a Δpgm , expression of avirulent nonpigmented phenotype (29) due to \sim 100-kb spontaneous chromosomal deletion (47). *b* Met⁺, methionine meiotroph (21).

pVK1 pPCP::Tn1 (insert in silent region) 33
pVK2 pPCP::Tn1 (insert in pst) 33

significance, if any, of this phenomenon. In this report we demonstrate that soluble and biologically active α -Pla can be extracted from *Escherichia coli* LE392/pPCP1 by autolysis or with 1.0 M NaCl, α -Pla undergoes spontaneous conversion to b-Pla and smaller forms, and complete Pla-dependent posttranslational hydrolysis of Yops (but not YadA) also occurs in *Y. pseudotuberculosis* transformed with pPCP1. These findings are discussed within the context of current models defining the translocation of Yops into host cells by docked yersiniae (5, 19).

MATERIALS AND METHODS

Bacteria. Salient properties of bacteria and plasmids used in this study are shown in Table 1. Transformation of *E. coli* LE392 with pPCP1 and derivatives of the latter containing Tn5 were previously described (58). *E. coli* LE392/pPCP1 was utilized in experiments concerned with extraction of soluble Pla and conversion of a-Pla to smaller derivatives. *E. coli* K802/pEK4 served as a source of Pla for generation of specific antiserum (6). DNA of pVK1 and pVK2 was isolated by the procedure of Clewell and Helinski (18) and transformed into *Y. pseudotuberculosis* strains 85 and 861 by freeze-thawing (20); transformants were selected on solid medium containing ampicillin. To minimize the potential of laboratory infection by Lcr⁺ cells of *Y. pseudotuberculosis* possessing pPCP1, experiments concerned with posttranslational hydrolysis of Pla and Yops in *Y. pseudotuberculosis* were undertaken with an avirulent *gua* isolate (intravenous 50% lethal dose $> 10^7$ bacteria) of strain PB1/+. The latter was induced and

selected by methods previously described for *Y. pestis* (7) and then coconjugated with F^+ to receive p2PCP::Tn1 containing a single copy of Tn1 in a silent region, as judged by mapping with restriction endonucleases (41). A methionine meiotroph (21) of nonpigmented (29) Lcr⁺ Pst⁺ *Y. pestis* KIM (substrain D39) and isogenic Lcr⁻ and Pst⁻ derivatives were used in comparative experiments. In mice Pgm⁻ isolates are virulent (50% lethal dose \approx 10 bacteria) when administered by intravenous injection but avirulent (50% lethal dose $> 10⁷$ bacteria) when administered by peripheral routes (30, 63) due to the mutational loss of an \sim 100-kb chromosomal sequence (23, 38) encoding the ability to absorb hemin, enzymes required for yersiniabactin synthesis, and the pesticin-yersiniabactin receptor ($\triangle pgm$) (47). Pst⁻ derivatives of *Y. pestis* were obtained by elimination of pPCP by cold-curing (56).

Retention of pPCP and its derivatives in yersiniae and *E. coli* was monitored in isolates recovered from experimental animals and in cultures used for experiments by determining the percentage of pesticinogenic colonies on pesticin agar, with *Y. pseudotuberculosis* PB1/0 used as an indicator (11). This value was always 100%, indicating that pPCP is retained tenaciously in *Y. pseudotuberculosis* and *E. coli*, as is known to be the case in *Y. pestis* (56).

Cultivation. Bacteria were grown at 37°C in liquid media contained in Erlenmeyer flasks (10%, vol/vol) aerated at 200 rpm on a water bath shaker (model G76; New Brunswick Scientific Co., New Brunswick, N.J.). *E. coli* K802/pEK4 was cultivated until late logarithmic growth phase in Luria-Bertani medium (Difco Laboratories, Inc., Detroit, Mich.) supplemented with 50 μ g of ampicillin (Sigma Chemical Co., St. Louis, Mo.) per ml. The culture then received 20 μ g of chloramphenicol (Sigma) per ml and was incubated for an additional 12 h at the same temperature before preparation of supernatant fluids. A minor modification of pesticin medium (28) used here for preparation and characterization of Pla consisted of 2.5% Sheffield NZ Amine (Kraft Inc., Memphis, Tenn.), 0.5%

FIG. 1. Silver stain (A) and immunoblots of unabsorbed (B) and absorbed (C) rabbit polyclonal anti-Pla following SDS-PAGE of purified Pla (lanes 1 and 12), whole cells of *Y. pestis* KIM cured of pPCP1 (substrain D45) (lane 2), *Y. pestis* KIM/pPCP1 (substrain D27) (lane 3), *E. coli* LE392 (lane 4), *E. coli* LE392/pPCP1 (lane 5), *E. coli* LE392/p2PCP::Tn1 lane 6). *Y. pseudotuberculosis* PB1/0 (lane 7), *Y. pseudotuberculosis* PB1/0/pPCP1 (substrain B59) (lane 8), *Y. pseudotuberculosis* PB1/0/p2PCP:Tn1 (lane 9), *Y. pseudotuberculosis* PB1/0/pVK1 (lane 10), and *Y. enterocolitica* WA (lane 11). Molecular mass markers (in kilodaltons) are provided in the left margins. The procedure used to absorb the antiserum is described in Materials and Methods. Note that only a-Pla was expressed by *E. coli*/pPCP1, whereas *Y. pseudotuberculosis* PB1/0/pPCP1 produced β -Pla as did *Y. pestis* KIM/pPCP1 (which also generated a third, ~31-kDa form, termed γ -Pla).

yeast extract (Difco), the salt component of Higuchi's defined medium (25) (final concentrations of 0.025 M K₂HPO₄, 0.01 M citric acid, 2.5 mM MgCl₂, 0.1 mM FeSO₄, and 0.01 mM MnCl₂), 25 mM potassium D-gluconate, and 2.5 mM sodium thiosulfate (all adjusted to pH 7.0 with 10 N NaOH). Pulse-chase experiments were performed with the defined medium of Zahorchak and Brubaker (68) supplemented with 0.5 mM guanosine and 0.5 mM hypoxanthine but lacking added L-methionine.

Antisera. To produce anti-Pla, cells of *E. coli* K802/pEK4 were grown as described above and removed by centrifugation $(10,000 \times g$ for 15 min), and the supernatant fluid was passed through a 0.2 - μ m-pore-size low-protein-binding membrane filter (Gelman Sciences, Ann Arbor, Mich.). Solid $(NH₄)₂SO₄$ was then added to the resulting sterile filtrate, and the 30 to 70% saturated fraction was collected by centrifugation (12,000 $\times g$ for 60 min), suspended in 0.025 M Tris \cdot Cl, pH 7.5, and dialyzed at 4°C against three changes of the same buffer. This material was applied in volumes of 2.0 ml containing 11.6 mg of protein to a column (1.6 by 60 cm) of Sephadex G-200 (Pharmacia, Uppsala, Sweden) equilibrated with 0.02 M Tris \cdot Cl buffer, pH 7.5, containing 0.2 M NaCl and 0.02% NaN₃, and then eluted at a flow rate of 0.4 ml/min in the same buffer. Eluted fractions (2.0 ml) were assayed for biological activity by use of fibrin plates as described previously (2). Positive samples were assayed by SDS-PAGE, pooled, and lyophilized. The resulting material was nearly homogeneous and, following immunization of rabbits, yielded antisera capable of recognizing Pla. To eliminate natural antibodies directed against *Yersinia* spp. and *E. coli*, the serum received disrupted and lyophilized cells of *E. coli* K-12, *Y. pseudotuberculosis* PB1, and *Y. enterocolitica* WA (20 mg apiece/ml) and was gently stirred for 30 min at 37°C and then overnight at 4°C followed by removal of particulate material by centrifugation (10,000 \times *g* for 30 min). After this cycle was repeated three times, gamma globulin was isolated by chromatography on DEAE cellulose as described previously (64).

Extraction of Pla. Bacteria were grown as noted above and harvested by centrifugation (10,000 \times *g* for 30 min), and the original culture supernatant fluid was retained for analysis by SDS-PAGE and immunoblotting. The organisms were washed once in 0.033 M potassium phosphate buffer, pH 7.0 (phosphate buffer), and then suspended at an optical density (620 nm) of 50 in either phosphate buffer alone or 1.0 M NaCl in phosphate buffer. These preparations, contained in 500-ml centrifuge bottles (50%, vol/vol), were vigorously shaken (250 rpm) on an Orbit platform shaker (model 3520; Lab-Line Instruments, Inc., Melrose Park, Ill.) for 30 min at 26°C and then centrifuged as described above. The resulting bacterial pellet was suspended in phosphate buffer and prepared for SDS-PAGE; supernatant fluids were similarly analyzed either directly or after passage through a 0.2-µm-pore-size low-protein-binding membrane filter (Gelman).

Pulse-chase determinations. The method of Sample et al. (57) was used to detect net synthesis and then degradation of Yops by Lcr^+ Pst⁺ yersiniae. Briefly, the technique involved restriction of vegetative growth at 37° C in Ca²⁺deficient medium lacking L-methionine and then pulsing the cultures for 15 s with carrier-free [³⁵S]methionine (New England Nuclear, Boston, Mass.) to yield a final concentration of 10 μ Ci/ml. The organisms were then chased by addition of excess unlabeled L-methionine to yield a final concentration of 1.6 mM. Samples of whole culture were removed at intervals, precipitated with an equal volume of cold 10% trichloroacetic acid, centrifuged $(5,000 \times g$ for 10 min), and prepared for SDS-PAGE. After staining, the gels were dried and exposed to X-ray film.

Miscellaneous. SDS-PAGE was performed by the method of Laemmli (34); separated proteins were visualized by silver staining (45). Alternatively, Coomassie blue was utilized for staining Pla; this procedure and that used for immunoblotting with the rabbit polyclonal anti-Pla described above have been described previously (9). SDS-PAGE of whole or processed bacteria was undertaken with \sim 10 µg of protein/lane; samples of extracted bacteria contained \sim 1 µg of protein. Protein quantity was estimated by the method of Lowry et al. (37). Outbred white mice (4 to 8 weeks) received bacteria by subcutaneous injection (10 mice per dose) to determine virulence. The 50% lethal dose was calculated by the method of Kerber as modified by Ashmarin and Vorob'iov (1). Appropriate measures were used for care and housing of laboratory animals.

RESULTS

Detection of Pla. Cells of *E. coli* and various yersiniae carrying pPCP1 were subjected to SDS-PAGE and then silver stained. Pla was not visible in the resulting gel (Fig. 1A), although bands corresponding in molecular mass to α -Pla (35) kDa) and β-Pla (33 kDa) plus numerous cross-reacting antigens were observed upon immunoblotting with unabsorbed rabbit antiserum raised against purified Pla (Fig. 1B). Immunoblots specific for Pla were obtained by absorption of this antiserum followed by purification of remaining immunoglobulin G (Fig. 1C). Only a-Pla was detected in *E. coli* LE392/ pPCP1, whereas both α - and β -Pla were produced by *Y*. *pseudotuberculosis* PB1/0/pPCP1. Cells of *Y. pestis* KIM/pPCP1 expressed both the α and β forms of Pla as well as a smaller variant of \sim 31 kDa, termed γ -Pla (Fig. 1C).

Solubilization of α **-Pla.** In no case was Pla expressed in culture supernatant fluids (Fig. 2, lanes 1). We attempted its controlled solubilization and defined its conversion to degradation products by extraction of Pst^+ bacteria with phosphate buffer alone or with 1.0 M NaCl in phosphate buffer. Only the a form was detected at significant levels in whole cells of *E. coli* LE392/pPCP1 after washing and suspension in phosphate buffer (Fig. 2A, lane 2). In contrast, similarly washed and

FIG. 2. Preparations obtained from *E. coli* LE392/pPCP1 (A), *Y. pseudotuberculosis* PB1/0/pPCP1 (substrain B59) (B), and *Y. pestis* KIM/pPCP1 (substrain D28) (C). Shown are culture supernatant fluid (lane 1); centrifuged whole cells suspended at an optical density (620 nm) of 50 in 0.033 M potassium phosphate buffer, pH 7.0 (phosphate buffer) (lane 2); supernatant recovered upon extraction (30 min) and then centrifugation of whole cells suspended in phosphate buffer (lane 3); supernatant recovered upon extraction (30 min) and then centrifugation of whole cells suspended in phosphate buffer after passage through a low-protein-binding membrane filter (lane 4); whole cells after extraction with phosphate buffer (lane 5); supernatant recovered upon extraction (30 min) and centrifugation of whole cells suspended in phosphate buffer plus 1.0 M NaCl (lane 6); supernatant recovered upon extraction (30 min) and centrifugation of whole cells suspended in phosphate buffer plus 1.0 M NaCl after passage through a low-protein-binding membrane filter (lane 7); and whole cells after extraction with phosphate buffer plus 1.0 M NaCl (lane 8). The positions of the α , β , γ , and δ forms of Pla are identified in the right margin.

suspended Lcr⁻ cells of *Y. pseudotuberculosis* PB1/0 (Fig. 2B, lane 2) and *Y. pestis* KIM harboring pPCP1 (Fig. 2C, lane 2) expressed α -Pla as well as the β and γ forms. Attempts to extract significant soluble Pla from the three organisms with phosphate buffer alone were not successful. In contrast, extraction with 1.0 M NaCl in phosphate buffer resulted in recovery from *E. coli* LE392/pPCP1 of α -, β -, and γ -Pla (Fig. 2A, lane 6). The same process did not release significant Pla in any form from Pst⁺ yersiniae (Fig. 2B, lane 6; Fig. 2C, lane 6), although it generated a small derivative in *Y. pestis*, termed δ-Pla, which remained associated with the bacteria (Fig. 2C, lane 8).

The α and β derivatives extracted from the surface of *E. coli* LE392/pPCP1 with 1.0 M NaCl in phosphate buffer were soluble, as demonstrated by their ability to pass through a lowprotein-binding filter, whereas smaller γ -Pla was lost during this procedure, indicating that it was denatured (Fig. 2A, lane 7). This observation illustrates that soluble Pla can be removed from *E. coli* LE392/pPCP1 (but not Pst⁺ yersiniae) by extraction with 1.0 NaCl in phosphate buffer. The filtered material obtained from *E. coli* LE392/pPCP1 with 1.0 M NaCl possessed potent biological activity ($>10^5$ U/10 μ l) as judged by direct assay on fibrin plates.

Pst⁺ phenotype of *Y. pseudotuberculosis.* A Pst⁺ isolate of a gua auxotroph of *Y. pseudotuberculosis* PB1/+ was selected by coconjugation of plasmid p2PCP::Tn1 accompanied by loss of F1. Comparison of this isolate with wild-type *Y. pestis* showed that both organisms expressed similar levels of pesticin, coagulase, and plasminogen activator activities. Subsequent mutational loss of pPCP1 in both species resulted in the disappearance of these activities (Table 2).

Following induction of the LCR, yersiniae were harvested, washed, and subjected to analysis by SDS-PAGE. After staining with silver (Fig. 3A) or Coomassie blue (Fig. 3B) or immunoblotting with anti-Pla (Fig. 3C), major Yops were recognized by comparison with isogenic mutants lacking pCD (pYV), pPCP, or both. Major discernible Yops in Lcr ⁺ Pst⁻ cells of *Y. pseudotuberculosis* (Fig. 3A and B, lanes 3) and *Y. pestis* (Fig. 3A and B, lanes 8) possessed molecular weights corresponding to values previously reported for YopH, YopC, YopD, and YopE (47). Those known to be expressed at lower concentrations were not detected by this procedure, nor were any Yops observed in Lcr⁺ Pst⁺ cells of *Y. pseudotuberculosis* (Fig. 3, lanes 1) or *Y. pestis* (Fig. 3, lanes 6). An Lcr^+ -specific

TABLE 2. Specific activities of various products in whole cells of *Y. pseudotuberculosis* PB1/+ and *Y. pestis* KIM

	Sp act (U/mg of protein) ^{<i>a</i>} of:			
Organism	pPCP	Pesticin	Coagulase	Plasminogen activator
Y. pestis	$+^b$	46,000	2.0	840
Y. pestis	θ	\cup	$_{0}$	
Y. pseudotuberculosis	$^{+}$	51,500	2.4	925
Y. pseudotuberculosis	θ	\cup		

^a Units are defined according to Straley and Brubaker (64).

^{*b*} +, carriage of pPCP.

FIG. 3. Effect of pPCP1 on expression of Yops in whole bacteria subjected to silver staining (A), Coomassie blue staining (B), and immunoblotting with rabbit polyclonal anti-Pla (C). Shown are the *Y. pseudotuberculosis* PB1/pPCP1, pYV (substrain B53) (lane 1); *Y. pseudotuberculosis* PB1/pPCP1 (substrain B59) (lane 2); *Y. pseudotuberculosis* PB1/pYV (substrain B44) (lane 3); *Y. pseudotuberculosis* PB1 (substrain B45) (lane 4); soluble (filtered) Pla extracted from *E. coli* LE392/pPCP1 in 1.0 M NaCl in 0.033 M potassium phosphate buffer, pH 7.0 (lane 5); *Y. pestis* KIM/pPCP1, pCD1 (substrain D27) (lane 6); *Y. pestis* KIM/pPCP1 (substrain D28) (lane 7); *Y. pestis* KIM/pCD1 (substrain D45) (lane 8); and *Y. pestis* KIM (substrain D47) (lane 9). (A and B) The solid arrows to the left of the gels indicate individual Yops, and the open arrows designate V antigen. (C) Arrows to the left of the immunoblot indicate the α , β , and γ forms of Pla.

peptide of \sim 37 kDa, established as V antigen, was observed in both Lcr ⁺ yersiniae (Fig. 3A and B, lanes 1, 3, 6, and 8). No form of Pla was detectable via silver staining (Fig. 3A, lane 5), but both α - and β -Pla were revealed by staining with Coomassie blue (Fig. 3B, lane 5) and by immunoblotting (Fig. 3C).

Pla-mediated degradation of Yops. A 15 s pulse with [³⁵S]methionine followed by a 1-h chase with excess unlabeled methionine was undertaken to demonstrate synthesis and concomitant degradation of Yops in Lcr⁺ Pst⁺ cells of *Y. pseudotuberculosis* PB1 (substrain B53) maintained in the LCR. Radioactive proteins in samples removed at intervals throughout this period were separated by SDS-PAGE and subjected to silver staining (Fig. 4A) and radiography (Fig. 4B). Lcr^+ -specific peptides of ~ 76 (YopF), 46 (YopH), 44 (YopB), 42 (YopC), 34 (YopD), and 25 (YopE) kDa underwent prompt posttranslational degradation and were not detected after 1 h. The stained gel (Fig. 4A, lanes 1 to 10) also showed the presence of YadA (\sim 200 kDa) and three similarly high molecular weight putative degradation products of this adhesin; native YadA was not well labeled within the short pulse used in these experiments (Fig. 4B, lanes 1 and 11).

A parallel determination with *Y. pestis* KIM (substrain D39) was performed for purposes of comparison (Fig. 5). One major difference was the anticipated absence of YadA and its presumed degradation products. The kinetics of hydrolysis of YopF, YopH, YopB, and YopC were similar between the two

species, whereas degradation of YopD occurred more slowly in Pst⁺ cells of *Y. pseudotuberculosis* (where its destruction required 1 h, as opposed to 8 min in *Y. pestis*). Both yersiniae converted YopE into its stabilized \sim 24-kDa degradation product. This process, however, was accomplished in 1 min by *Y. pestis*, as opposed to 15 min for *Y. pseudotuberculosis*.

Virulence of Pst⁺ derivatives of *Y. pseudotuberculosis***. In** order to determine if expression of Pla by *Y. pseudotuberculosis* influences virulence in mice, we compared the subcutaneous 50% lethal doses of wild-type isolates transformed with Pst plasmids (pVK1 and pVK2). As shown in Table 3, carriage of these plasmids did not significantly enhance invasion of tissues as judged by only modest decreases in virulence.

DISCUSSION

Pla possesses significant homology to the outer membrane protease OmpT of *E. coli* (59) now known to be active under extreme denaturing conditions (66). OmpT from certain clinical isolates of *E. coli* also activated plasminogen, although additional features of surface architecture unique to pathogenic strains may have contributed to this capability (39). It is therefore of interest that the N terminus of native OmpP (a similar outer membrane protease possessing 87% identity to OmpT) was vulnerable to proteolytic digestion whereas its C-terminal region was protected. This finding indicated that

FIG. 4. Silver-stained gel (A) and corresponding autoradiogram (B) of trichloroacetic acid-precipitated material from cultures of *Y. pseudotuberculosis* PB1 cultivated at 37°C in Ca²⁺-deficient medium. Lcr⁺ Pst⁺ ce for 0 s (lane 1), 15 s (lane 2), 30 s (lane 3), 1 min (lane 4), 2 min (lane 5), 4 min (lane 6), 8 min (lane 7), 15 min (lane 8), 30 min (lane 9), and 1 h (lane 10). Control lanes (pulsed for 15 s and then chased for 1 h) contained isogenic Lcr⁺ Pst⁻ (substrain B44) (lane 11), Lcr⁻ Pst⁺ (substrain B59) (lane 12), and Lcr⁻ Pst⁻ (substrain B59) (lane 12), and Lcr⁻ Pst⁻ (substrain Bla and p24 indicate b-lactamase and the stable YopE degradation product (11), respectively. Note the presence of YadA and its derivatives in the silver stain.

the catalytic center of OmpP is located at or near the N terminus and that its C-terminal end promotes attachment to the outer membrane (32). Similarly, introduction of hexahistidine at the N terminus of Pla prevented biological activity whereas its addition at the C terminus prevented anchoring (unpublished observations). In view of these findings, we abandoned the prospect of engineering a soluble form of Pla and considered alternatives known to facilitate physical removal of native membrane proteins, including autolysis and extraction with monovalent cations (14, 36, 53).

These processes failed to release Pla in active form from *Y. pestis* or *Y. pseudotuberculosis* carrying pPCP, although particulate samples of high specific activity were sometimes obtained. However, their application to certain isolates of *E. coli* transformed with pPCP, including strains K802 and LE392, resulted in extraction of active enzyme. The reason for this difference is unknown, but a likely possibility is that Pla is processed and inserted into the outer membrane of *E. coli* LE392 exactly like OmpT or OmpP but that differences in primary structure result in faulty anchoring and thus permit solubilization upon autolysis or increased ionic strength. We emphasize that this phenomenon was observed only for the noted *E. coli* K-12 derivatives and does not apply to all strains of the species.

FIG. 5. Silver-stained gel (A) and corresponding autoradiogram (B) of trichloroacetic acid-precipitated material from cultures of *Y. pestis* KIM (substrain D39) and corresponding derivatives. See the legend to Fig. 4 for details.

Although the site cleaved by plasminogen activators to generate plasmin is established (55), essentially nothing is known about the posttranslational conversion of Pla, the mechanism whereby Pla degrades Yops, or the biological significance of this reaction. Inability to define these processes reflects the fact that heretofore Pla could not be prepared in a form sufficiently homogeneous to rigorously determine its mechanisms of conversion and catalysis. These issues can now be addressed by using the rabbit polyclonal antiserum raised against Pla from *E. coli* LE392/pPCP1.

Of initial interest was the finding that cells of this isolate grown in an enriched medium (28) expressed only cell-bound a-Pla as opposed to those of *Y. pseudotuberculosis* PB1/0/ pPCP1 or *Y. pestis* KIM/pPCP1, which also formed β-Pla. These results are in contrast to those of Sodeinde and Goguen

TABLE 3. The influence of pPCP on the subcutaneous 50% lethal dose of Lcr⁺ cells of *Y. pseudotuberculosis* in mice

Strain	pPCP	50% lethal dose	\mathbb{R}^d
85		6.7×10^{1}	
85/1	$+$ (pVK1) ^b	8.2×10^{2}	12.2
85/2	$+$ (pVK2)	2.1×10^3	31.3
861		7.5×10^{5}	
861/1	$+$ (pVK1)	6.2×10^{6}	8.3
861/2	$+$ (pVK2)	4.8×10^{6}	6.4

^a R, ratio of 50% lethal dose of the parent to that of the isogenic derivative transformed with pPCP.

 b +, carriage of pPCP.

(58), who detected cell-bound α - and β -Pla in *E. coli* cultivated in a minimal medium. We have repeated this observation with *E. coli* LE392/pPCP1 but have not yet identified variables that account for the appearance of β -Pla under austere conditions. However, we noted that the process of extracting *E. coli* LE392/pPCP1 with NaCl after growth in the enriched medium generated both β - and γ -Pla, suggesting that the smaller forms arise as sheared derivatives of α -Pla. Extraction failed to remove soluble forms of Pla from Pst⁺ cells of *Y. pestis* or *Y. pseudotuberculosis* (but did generate cell-bound γ - and δ -Pla), thus emphasizing that Pla is tenaciously bound to the outer membrane of yersiniae.

The anti-Pla serum enabled precise localization of α -Pla and its derivatives in stained gels. These determinations prompted the unexpected observation that Pla cannot be visualized by silver staining. Fortunately, Pla stained readily with Coomassie blue, which we recommend for determining its presence by SDS-PAGE. These experiments provided assurance that *Y. pseudotuberculosis* PB1/+/pPCP1 does indeed express levels of Pla similar to those in wild-type *Y. pestis*, thereby permitting a direct comparison of Yop degradation by the two isolates.

Yops of Lcr⁺ Pst⁺ cells of *Y. pseudotuberculosis* underwent hydrolysis in a manner similar to that previously reported for *Y. pestis*. Two distinctions, however, were noted between the two species. First, at least portions of the process occurred more rapidly in *Y. pestis*, even though the specific activities of Pla in the two isolates were equivalent as determined by fibrinolytic assay. This phenomenon is illustrated by the prompt disappearance of YopE in Pst⁺ cells of *Y. pestis* KIM with immediate conversion to its stable derivative (13), as opposed to more leisurely degradation in Lcr⁺ Pst⁺ cells of *Y. pseudotuberculosis* PB1. Second, YadA of the latter underwent evident Plamediated alteration, resulting in the appearance of four distinct structural elements, with the molecular mass of the largest equal to that of native YadA in the Lcr ⁺ Pst⁻ control. The significance of this observation is not fully understood, but these components probably represent native YadA and the YadA heteropolymeric subunits described by Gripenberg-Lerche et al. (24).

No major differences in virulence were found between wildtype *Y. pseudotuberculosis* and its Pst⁺ derivatives. Accordingly, expression of Pla did not significantly increase the virulence of these strains of *Y. pseudotuberculosis* by subcutaneous injection as is known to occur in *Y. pestis* (8). This difference emphasizes that avirulence caused by mutational loss of a native plasmid does not guarantee that acquisition of that plasmid by a naive bacterium will necessarily assure enhanced virulence. Many possibilities could account for this distinction, including pPCP-mediated disruption of pYV partitioning, interference with iron assimilation by pesticin, and hydrolysis of some unique *Y. pseudotuberculosis*-specific virulence factor by Pla. At present, we favor the notion that YadA, known to reduce the virulence of *Y. pestis* (54), remains at least partially intact in *Y. pseudotuberculosis* harboring pPCP (and thus may still retard dissemination of the bacteria from peripheral sites of injection). Resolution of this inconsistency is beyond the scope of the present study.

Major pCD (pYV)-encoded virulence factors include the cytotoxic Yops noted previously. Despite the essential nature of these proteins, they are promptly degraded by Pla during expression in vitro. Recent work on the nature of the LCR and results reported here may provide a basis for understanding this enigma. Classical studies on the regulation of macromolecular synthesis in prokaryotes have established the existence of global regulatory mechanisms that assure constant ratios of protein to DNA, thereby preventing lethal unbalanced growth

(40). Bacteriostasis of yersiniae in vitro resulting from $Ca²⁺$ privation at 37°C reflects termination of ongoing rounds of DNA synthesis but continued synthesis of mRNA (15, 68, 69) known to largely encode pCD encoded functions, especially Yops (42, 43). The ratio of DNA to protein would thus become skewed if Yops were permitted to accumulate within the cytoplasm of static Ca^{2+} -starved bacteria; therefore, a process is required to assure their elimination, either by turnover or export into culture supernatant fluid. Both mechanisms are utilized as judged by the occurrence of type III secretion and concomitant Pla-mediated degradation. It is not yet clear if an identical secretion process also accounts for translocation of Yops into host cell cytoplasm by docked yersiniae. In this context, it may be significant that YopE can be secreted by two distinct systems (16); perhaps only one such mechanism is linked to subsequent Pla-mediated hydrolysis.

In any event, it is now evident that cytotoxic Yops are innocuous to the host unless translocated during bacterium-host cell contact (19). Accordingly, the ability of Pla to hydrolyze excreted but not translocated Yops is unlikely to interfere with expression of virulence. Indeed, this function might even favor the ability of *Y. pestis* to cause acute disease by eliminating soluble Yops from Ca^{2+} -deficient necrotic lesions in visceral organs where they would otherwise be recognized as foreign proteins. It is significant in this regard that visceral lesions in mice formed by Lcr⁺ cells of *Y. pseudotuberculosis* arose as abscesses surrounded by professional phagocytes, whereas those generated by *Y. pestis* appeared as pure necrotic foci entirely lacking in signs of inflammation (46, 65).

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