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Accessibility
Poly-N-Acetylg glucosamine Expression by Wild-Type Yersinia pestis Is Maximal at Mammalian, Not Flea, Temperatures

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ABSTRACT Numerous bacteria, including Yersinia pestis, express the poly-N-acetylg glucosamine (PNAG) surface carbohydrate, a major component of biofilms often associated with a specific appearance of colonies on Congo red agar. Biofilm formation and PNAG synthesis by Y. pestis have been reported to be maximal at 21 to 28°C or “flea temperatures,” facilitating the regurgitation of Y. pestis into a mammalian host during feeding, but production is diminished at 37°C and thus presumed to be decreased during mammalian infection. Most studies of PNAG expression and biofilm formation by Y. pestis have used a low-virulence derivative of strain KIM, designated KIM6+, that lacks the pCD1 virulence plasmid, and an isogenic mutant without the pigmentation locus, which contains the hemin storage genes that encode PNAG biosynthetic proteins. Using confocal microscopy, fluorescence-activated cell sorter analysis and growth on Congo red agar, we confirmed prior findings regarding PNAG production with the KIM6+ strain. However, we found that fully virulent wild-type (WT) strains KIM and CO92 had maximal PNAG expression at 37°C, with lower PNAG production at 28°C both in broth medium and on Congo red agar plates. Notably, the typical dark colony morphology appearing on Congo red agar was maintained at 28°C, indicating that this phenotype is not associated with PNAG expression in WT Y. pestis. Extracts of WT sylvatic Y. pestis strains from the Russian Federation confirmed the maximal expression of PNAG at 37°C. PNAG production by WT Y. pestis is maximal at mammalian and not insect vector temperatures, suggesting that this factor may have a role during mammalian infection.

IMPORTANCE Yersinia pestis transitions from low-temperature residence and replication in insect vectors to higher-temperature replication in mammalian hosts. Prior findings based primarily on an avirulent derivative of WT (wild-type) KIM, named KIM6+, showed that biofilm formation associated with synthesis of poly-N-acetylg glucosamine (PNAG) is maximal at 21 to 28°C and decreased at 37°C. Biofilm formation was purported to facilitate the transmission of Y. pestis from fleas to mammals while having little importance in mammalian infection. Here we found that for WT strains KIM and CO92, maximal PNAG production occurs at 37°C, indicating that temperature regulation of PNAG production in WT Y. pestis is not mimicked by strain KIM6+. Additionally, we found that Congo red binding does not always correlate with PNAG production, despite its widespread use as an indicator of biofilm production. Taken together, the findings show that a role for PNAG in WT Y. pestis infection should not be disregarded and warrants further study.
and succinylated wheat germ agglutinin, both of which have binding specificity for β-GlcNAc moieties (11). Taken together, these findings strongly indicate that *Y. pestis* produces a surface polysaccharide composed of PNAG.

Thus far, the major pathogenic role of the *hms*-dependent biofilms formed by *Y. pestis* is in the colonization of the proventricular structure of fleas, a crucial step in the transmission of plague bacteria (1, 3). That conclusion was strengthened by the belief that the *Y. pestis* biofilm was maximally formed at a temperature of ~28°C but poorly or not at all at mammalian temperatures, i.e., ≥37°C (12). However, a review of the papers reporting this attribute of biofilm formation has revealed that this finding is based on results obtained with a single avirulent derivative of *Y. pestis* strain KIM, the *hms*<sup>+</sup> strain designated KIM6<sup>+</sup> by Perry and coworkers, and an isogenic variant designated KIM6, which lacks the 102-kb *pgm* locus that contains the *hms*<sup>HEFS</sup> operon and which is referred to here as KIM6 (Δ*pgm*) (2–4, 8, 12–16). KIM6<sup>+</sup> was derived from wild-type (WT) strain KIM by deletion of the pCD1 virulence plasmid which contains the genes for the type III secretion proteins that are the major determinants of high-level virulence in plague bacteria. Erickson et al. (9) found that when they cloned a PNAG-degrading β-hexosaminidase from *Y. pseudotuberculosis* into KIM6<sup>+</sup> *Y. pestis* there was reduced biofilm formation *in vitro* and a reduction in the proventricular blockage of fleas. However, microarray analysis of WT *Y. pestis* recovered from the lungs of infected mice showed that genes of the *hms* operon were upregulated during experimental pneumonic infection (17, 18), although its role in active infection has not been delineated. Lack of data on PNAG expression by WT *Y. pestis* and increased transcription of the *hms* genes during experimental mouse infection with WT strains suggested the possibility that WT *Y. pestis* does not show the same variation in PNAG production at 28 and 37°C as the KIM6<sup>+</sup> derivative does. We thus investigated PNAG production among WT *Y. pestis* strains KIM and CO92 and sylvatic strains from the Russian Federation, where plague is endemic, to determine if they regulate polysaccharide synthesis in the same temperature-dependent manner as strain KIM6<sup>+</sup>.

**RESULTS**

Detection of PNAG by confocal microscopy and FACS analysis of WT and pCD1-negative strains of *Y. pestis* grown at 28 or 37°C. Using *Y. pestis* cells grown in brain heart infusion broth (BHIB), we confirmed results from prior publications (12, 13, 15, 16) showing the pCD1-negative KIM6<sup>+</sup> strain of *Y. pestis* exhibited strong production of PNAG at 28°C, as evidenced by confocal immunofluorescence microscopy (Fig. 1A), which was markedly diminished at 37°C. The KIM6 (Δ*pgm*) strain, without the *pgm* locus, which also lacks the *hms* biosynthetic genes for PNAG, showed the same lack of reactivity with Alexa Fluor 488 (AF488)–conjugated monoclonal antibody (MAb) F598 to PNAG as it did with control MAb F429-AF488 specific to *P. aeruginosa* alginate (Fig. 1A). Treatment of KIM6<sup>+</sup> cells grown in BHIB at 28°C with chitinase, which has no effect on PNAG, retained reactivity with MAB F598-AF488, whereas digestion with the PNAG-hydrolyzing enzyme dispersin B markedly diminished the reactivity of cells with MAB F598-AF488 (Fig. 1B).

In contrast to the data contained in numerous reports on the use of these two KIM6 variants (12, 13, 15, 16), we found that PNAG expression on cells of pCD1-positive WT *Y. pestis* KIM grown in BHIB at 37°C was greater than that on cells grown at 28°C (Fig. 2A). Similarly, WT *Y. pestis* CO92 cells showed little to no PNAG expression at 28°C in BHIB and marked expression at 37°C (Fig. 2A), which is the complete opposite of the results obtained over the past decades with the pCD1-negative derivatives of strain KIM (KIM6+). These outcomes were obtained with three separate samples of WT cells provided by the NERCE (New England Research Center for Excellence) Microbiology and Animal Core facility, WT *Y. pestis* KIM or CO92 cells grown in BHIB and treated with chitinase retained reactivity with MAB F598-AF488, whereas dispersin B treatment resulted in loss of immunoreactivity (Fig. 2B), indicative of PNAG production by WT *Y. pestis* cells.

Fluorescence-activated cell sorter (FACS) analysis of PNAG expression by the KIM pCD1-negative *Y. pestis* variants and WT strains KIM and CO92 following growth in BHIB correlated with the confocal immunofluorescence microscopy findings (Fig. 3A). For the pCD1-negative KIM6+ strain, FACS analysis showed higher levels of PNAG production when cells were grown at 28°C, with little made by cells grown at 37°C, while the *hms*-negative KIM6 strain had no reactivity above that of the control when grown at either 28 or 37°C (Fig. 3A, top). For WT strains KIM and CO92, FACS confirmed the higher expression of PNAG by cells grown at 37°C than by those grown at 28°C (Fig. 3A, bottom). Quantification of the mean fluorescence intensity obtained when comparing WT *Y. pestis* cells grown at either 28 or 37°C using MAB F598-AF488 to PNAG indicated that strain KIM produced about 2.6 times as much PNAG as with control MAB F498-AF488 at 28°C, while at 37°C, the expression of PNAG increased to 14 times the background level and was comparable to that of the
pCD1-negative KIM6+ strain grown at 28°C (Fig. 3B). For WT Y. pestis strain CO92, the FACS analysis also validated the confocal microscopic analysis in that only a small amount of PNAG was detected in the same region of the world, whereas strain 1146 produced PNAG in a manner analogous to that of WT strain KIM, which also originated in the same region of the world, whereas strain 1146 produced PNAG in a manner analogous to that of WT Y. pestis strain CO92, with little detected at 28°C and more detected at 37°C. None of these isolates behaved like the pCD1-negative KIM6+ strain, which showed greater PNAG production at 28°C than at 37°C.

**DISCUSSION**

The **hms** operon of *Y. pestis* strain KIM6+ is required for biofilm formation *in vitro* and the production of an aggregate of *Y. pestis* cells in the flea foregut (9), phenotypes linked to the transmission of this pathogen from this insect vector to a mammalian host (1, 3). The initial phenotype associated with the expression of the **hms** genes in *Y. pestis* was production of dark red colonies on Congo red agar plates (12), and this was later linked to PNAG production and biofilm formation (8, 9). Our findings do not dispute these previously published reports indicating that PNAG is synthesized maximally by *Y. pestis* KIM6+ at 28°C, as we also found higher PNAG production by this strain at 28°C, as determined by confocal microscopy and FACS analysis. However, it now appears that synthesis of PNAG by WT *Y. pestis* strains is not associated with the dark colony phenotype on Congo red agar, as we found that this phenotype occurred at 28°C in all of the *Y. pestis* stains tested, whereas PNAG production by WT strains was maximal at 37°C. Thus, the reported temperature regulation that links the maximal expression at 28°C of the phenotypes of dark red colonies on Congo red agar, PNAG production, and biofilm formation does not appear to occur in WT *Y. pestis*. Congo red is a somewhat
nonspecific dye that has been shown to bind multiple polysaccharides (20), as well as proteins (21). Its reactivity with *Y. pestis* strains at lower temperatures suggests a possible interaction with a non-PNAG temperature-regulated molecule.

Given our findings that the KIM6+ strain does not regulate PNAG production as do WT *Y. pestis* strains (including the WT KIM strain from which KIM6+ was derived), PNAG production by KIM6+ may not be representative of what would occur in a comparison of WT *Y. pestis* with an isogenic Δ*hms* strain. It is likely that some genetic event occurred in the KIM6+ strain leading to the differential regulation of *hms* genes and/or Hms proteins that is not representative of WT *Y. pestis*. This could be attributed to the fluidity of the *Y. pestis* genome, which contains a large number of insertion sequences, predisposing it to chromosomal deletions and rearrangements (22, 23).

In *Y. pestis*, biofilm formation and dispersion are largely regulated by two proteins, HmsT and HmsP (14). These proteins bridge the signal cascade between transmembrane sensors that detect environmental changes and downstream proteins involved in changing bacterial physiology in response to that signal, using cyclic di-GMP as the intermediate messenger (24, 25). HmsT is a diguanylate cyclase that synthesizes cyclic di-GMP, while HmsP is a phosphodiesterase that degrades cyclic di-GMP (26, 27). In KIM6+, several Hms proteins are posttranslationally regulated by protease degradation upon a temperature increase from 26 to 37°C (12, 14). Degradation of HmsT, the diguanylate cyclase that triggers biofilm formation at 37°C (in KIM6+) could explain the biofilm reduction seen at that temperature. While we have not attempted to pinpoint the differences between KIM6+ and WT *Y. pestis* strains that could have resulted in differential PNAG production, it seems plausible to hypothesize that differences or mutations in any protein(s) involved in the complex signal cascade could change the amount of PNAG synthesized in response to a temperature shift. Additionally, quorum sensing in *Y. pestis* is also

FIG 3 (A) Expression of PNAG by the indicated strains of *Y. pestis* grown in BHIB at the temperatures indicated in individual panels as determined by FACS. Dark gray trace, reactivity with control MAb F429 to *P. aeruginosa* alginate. Light gray trace, reactivity with PNAG-specific MAb F598. (B) Quantitative analysis of fluorescence intensity of *Y. pestis* strains reacted with either control MAb F429 (black bars) or MAb F598 to PNAG (gray bars). Bars indicate means, and error bars indicate the standard errors of the means.

FIG 4 Colony morphology on Congo red agar of four *Y. pestis* strains grown overnight at the temperatures indicated. Only *pgm/hms*-positive strains produced dark-staining colonies at 28°C and not at 37°C.
temperature regulated, with a change in the protein expression profile when a high cell density is reached (28). Proteins involved directly in, or upstream of, PNAG production may be regulated differently in KIM6+ in response to the achievement of quorum density, potentially also resulting in changes in biofilm formation.

Overall, we have confirmed that Y. pestis produces PNAG antigen but have also shown that the avirulent KIM6+ derivative does not regulate PNAG production as do WT Y. pestis strains. We believe that the prior general conclusion of increased PNAG production and biofilm formation at 21 to 28°C by Y. pestis (12, 13) was due to the sole use of the KIM6+ strain to determine these phenotypes, whereas maximal PNAG production by WT Y. pestis occurs at 37°C. However, despite the increased production of PNAG at mammalian temperatures by WT Y. pestis, any role it may have in virulence has yet to be elucidated. Unfortunately, conclusions regarding the lack of a role for PNAG in the virulence of Y. pestis are confounded by the use of KIM6+ restored to full virulence by the reintroduction of pCD1 (15, 26). Strains lacking pgm are attenuated, although the reduction in virulence has been attributed largely to the loss of an iron uptake protein (Irp2) found within the pgm pathogenicity island (29, 30). Notably, pgm⁺Δirp2 strains are still more virulent than Δpgm strains, suggesting that other genes in the pgm genetic element play roles in infection and pathogenesis (31). In a study with a CO92 derivative, an inactivating mutation in hmsP (resulting in a strain deficient in biofilm breakdown) rendered the strain 100-fold more resistant than its parental strain to the antibiotic gentamicin (32). Strains lacking hmsP are attenuated, whereas maximal PNAG production by WT Y. pestis occurs at 37°C. However, despite the increased production of PNAG at mammalian temperatures by WT Y. pestis, any role it may have in virulence has yet to be elucidated. Unfortunately, conclusions regarding the lack of a role for PNAG in the virulence of Y. pestis are confounded by the use of KIM6+ restored to full virulence by the reintroduction of pCD1 (15, 26). Strains lacking pgm are attenuated, although the reduction in virulence has been attributed largely to the loss of an iron uptake protein (Irp2) found within the pgm pathogenicity island (29, 30). Notably, pgm⁺Δirp2 strains are still more virulent than Δpgm strains, suggesting that other genes in the pgm genetic element play roles in infection and pathogenesis (31). In a study with a CO92 derivative, an inactivating mutation in hmsP (resulting in a strain deficient in biofilm breakdown) rendered the strain 100-fold more resistant than its parental strain to the antibiotic gentamicin (32).

In the same study, an hmsP mutant was found to be deficient in the invasion of human lung fibroblast cells. Interestingly, ΔhmsP mutants were attenuated in mouse infection models, albeit these infections were established with KIM6+-derived strains (26). Taken together, these results suggest that while Hms proteins (and PNAG production) may be dispensable in the initial stage of infection, PNAG production could possibly confer an advantage on the bacteria once infection has been established. The finding that transcription of the hms genes by WT strain CO92 is increased during active pneumonic plague infection in a murine model (17, 18) would support the possibility that PNAG production and biofilm formation play a role later in infection, potentially protecting bacteria from antibiotics and host defenses. The fact that the KIM6+ derivative does not regulate PNAG production in vivo in the same way as WT Y. pestis does, coupled with the finding that PNAG production is maximal at 37°C in vitro and in a model of active plague infection, leaves open the possibility that PNAG may have a role in Y. pestis virulence.

**MATERIALS AND METHODS**

**Bacterial strains and growth conditions.** Y. pestis WT strains KIM and CO92, along with strains KIM6+ (pgm⁺ ΔhmsP), and KIM (Δpgm ΔhmsP⁺), kindly provided by Robert Perry, Louisville, KY, were used in these studies. All strains were verified as having the proper phenotype by the Microbiology and Animal Resources Core Laboratory of the NERCE in Biodefense and Emerging Infectious Diseases. For confocal microscopy and FACS analysis, all strains were grown at NERCE. A loopful of a frozen seed stock was streaked onto brain heart infusion agar and incubated for 2 days at 28°C. Next, several colonies were inoculated into 1 ml BHIB and mixed for 1 min and then 4 ml additional BHIB was added. The suspension was adjusted to an optical density at 650 nm of 1.0, and 1 ml thereof was inoculated into 50 ml of BHIB in a 250-ml vented, baffled flask and incubated at either 37 or 28°C with shaking at 200 rpm for 16 to 18 h. A similar procedure was used to grow colonies on Congo red agar plates for 48 h at either 28 or 37°C. The Congo red agar medium used (20 ml per plate) consisted of 37 g/liter of BHIB (Oxoid), 50 g/liter sucrose, 10 g/liter agar, and 400 mg/liter of sterile Congo red, which was added after sterilization by autoclaving, followed by cooling to 55°C.

After growth, cells were recovered by centrifugation, resuspended in 1 ml phosphate-buffered saline (PBS), and mixed thoroughly. Additional PBS was added (4 ml), and the cells were centrifuged again and resuspended in 1 ml of sterile 10% formalin. An additional 4 ml of 10% formalin was added, and the cultures were stored at 28°C for 4 days. Prior to the
addition of formalin, an aliquot was taken to ascertain the number of CFU/ml by dilution and plating on blood agar plates (tryptic soy agar plates supplemented with 5% sheep blood). The colony counts of all cultures grown at 28 or 37°C were between 3 × 10^8 and 4 × 10^8 CFU/ml. The sterility of the formalin-treated cultures was ascertained by adding 100 μl of each culture to duplicate tubes of BHIB that were then incubated for 4 days at 37°C. No growth in any sample was seen after formalin fixation. Samples were adjusted to a concentration of 3 × 10^8 CFU/ml, aliquoted, and frozen at −80°C until used.

Antibody to PNAG. Fully human MAb F598, which binds to both native and deacetylated glycoforms of the S. aureus PNAG molecule, was used in this study (33). Immune serum from a rabbit immunized with dPNAG-DT (34) was used for detection of PNAG by immunoblotting.

Confocal microscopy. For confocal microscopic visualization of PNAG production by Y. pestis, a 100-μl frozen aliquot of formalin-fixed cells was defrosted and centrifuged to recover the bacterial cells, which were suspended in 20 μl of PBS and then air dried on microscope slides. Samples were fixed to the slides by exposure to 100% methanol for 1 min at room temperature, washed twice with PBS, and then labeled either with a control human IgG1 MAb, F429, specific to P. aeruginosa alginate (35) or with PNAG-specific human IgG1 MAb F598 (33), both of which were directly conjugated to AF488 (final MAb-fluorophore concentration, 1 μg/μl).

![Confocal microscopy images](image)

**FIG 7** (A) Expression of PNAG by the indicated strains of Y. pestis grown on Congo red agar at the temperatures indicated in individual panels as determined by FACS analysis. Dark gray trace, reactivity with control MAb F429 to P. aeruginosa alginate. Light gray trace, reactivity with PNAG-specific MAb F398. (B) Quantitative analysis of fluorescence intensity of Y. pestis strains reacted with either control MAb F429 (black bars) or MAb F598 to PNAG (gray bars). Bars indicate means, and error bars indicate the standard errors of the means.

### TABLE 1 Summary of PNAG immunoblot assay results of cell surface extracts of diverse Russian laboratory or field isolates of Y. pestis

<table>
<thead>
<tr>
<th>Strain</th>
<th>Biovar/subspecies</th>
<th>pgm</th>
<th>PNAG antibody binding(^c) when grown at:</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>28°C</td>
</tr>
<tr>
<td>KIMD1</td>
<td>Medievalis/pestis</td>
<td>−</td>
<td>−</td>
</tr>
<tr>
<td>260(11)</td>
<td>Antigua/pestis</td>
<td>−</td>
<td>−</td>
</tr>
<tr>
<td>EV11M</td>
<td>Antigua/pestis</td>
<td>−</td>
<td>−</td>
</tr>
<tr>
<td>K1(780)(^a)</td>
<td>Medievalis/pestis</td>
<td>−</td>
<td>−</td>
</tr>
<tr>
<td>KM239</td>
<td>Medievalis/pestis</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>625(44)(^a)</td>
<td>Medievalis/pestis</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>260(5)</td>
<td>Antigua/pestis</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>U2357</td>
<td>Medievalis/altaica</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>P-1680(^a,b)</td>
<td>Antigua/caucasica</td>
<td>+</td>
<td>−</td>
</tr>
<tr>
<td>1146(^b)</td>
<td>Antigua/caucasica</td>
<td>+</td>
<td>−</td>
</tr>
</tbody>
</table>

\(^a\) This strain carries the pCD1 virulence plasmid.
\(^b\) This strain was not virulent in a guinea pig infection model.
\(^c\) −, no binding; +, positive binding; ++, exceptionally strong binding.
5.2 µg/ml) along with the DNA-binding dye Syto 83 at 4 µM in PBS containing 0.5% bovine serum albumin (BSA). Staining reaction mixtures were incubated overnight at 4°C. Slides were washed twice with PBS and mounted for visualization by confocal microscopy.

To verify that the material reacting with PNA-EC-specific MAB F598 was likely PNA-EC, additional formalin-fixed samples were washed and resuspended in 200 µl of Tris-buffered saline (pH 6.8) containing either 50 µg/ml of the PNA-EC-hydrolyzing enzyme dispersin B (36) (Kane Biotech, Manitoba, Canada) or the chitin-degrading enzyme chitinase (Sigma), which has no effect on PNA-EC, and incubated overnight at 37°C. Cells were washed and reacted with MAB F598-AF488 as described above. Retention of MAB F598- AF488 binding following digestion with chitinase with loss of binding following digestion with dispersin B is indicative that the target antigen is PNA-EC.

**FACS analysis**. Frozen aliquots of formalin-fixed samples of *Y. pestis* cells were thawed and vortexed, and 300 µl of each sample was used to recover cells by centrifugation. The cells were washed with PBS with 0.5% BSA and split into three 100-µl aliquots. Samples were incubated overnight at 4°C with either 100 µl of PBS-BSA (unlabeled, used for calibration) or 5.2 µg/ml of either MAB F429 or MAB F598 directly conjugated with AF488. Samples were washed twice with PBS and resuspended in 500 µl of PBS before filtering through filter-cap FACS tubes and analyzed on a BD FACS Calibur.

**Immunoblot assays**. Cell surface extracts obtained from *Y. pestis* strains from regions of the Russian Federation where plague is endemic in rodents (19) were subjected to treatment with protease K at 65°C for 5 h, followed by protease inactivation at 80°C for 30 min, and then spotted onto a nylon membrane that was provided to us by Andre Anisimov and colleagues at the State Research Center for Applied Microbiology, Obolensk, Moscow Region, Russian Federation. The membrane containing the bacterial cell surface extracts was incubated with antisera from rabbits immunized with dPNAG conjugated with DT (Sigma), which has no effect on PNAG, and incubated overnight at 37°C. Cells were washed and reacted with MAB F598-AF488 as described above. Depolymerization of β-1,6-N-acetyl-d-glucosamine during the emergence of *Yersinia pestis*. J. Bacteriol. 190:8163–8170.


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