Differences in growth characteristics and elementary body associated cytotoxicity between Chlamydia trachomatis ocuugenital serovars D and H and Chlamydia muridarum

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Aim: In vitro growth and elementary body (EB) associated cytotoxicity of two Chlamydia trachomatis strains belonging to serovars D and H and C muridarum were compared to identify differences that correlate with virulence variations between these strains in the mouse model of female genital tract infection, and phenotypic characteristics that could explain human epidemiological data on serovar prevalence and levels of shedding during serovar D and H infection.

Methods: Replication cycle kinetics, inclusion characteristics, and EB associated cytotoxicity were assessed in McCoy cell monolayers using culture, light microscopy, and lactate dehydrogenase release.

Results: Over 72 hours, more rapid production and release of inclusion forming units (ifu) allowed C muridarum to initiate two replication rounds, resulting in 4–8 times more ifu/unit of infection than with serovars D and H. Although C muridarum EBs were significantly more cytotoxic to McCoy cell monolayers than serovar D at moderate and high multiplicity of infection ratios (MOI), serovar H EBs were significantly more cytotoxic than C muridarum, even at the lowest MOI tested.

Conclusions: These phenotypic differences are consistent with the more invasive course and severe pathological outcome of infection in mice infected with C muridarum, providing an objective basis for questioning the appropriateness of C muridarum as a surrogate for the human biovar of C trachomatis in the murine model of female genital tract infection. The differences seen between the human strains could help explain human epidemiological data relating to differences in prevalence and level of shedding that occurs during infection with ocuugenital serovars D and H.

There are 19 distinct serotypes (serovars) of Chlamydia trachomatis that infect humans, with serovars D, Da, E, F, G, Ga, H, I, Ia, J, and K belonging to the group referred to as ocugential serovars, which are collectively the most common sexually transmitted bacterial agents throughout the world (chlamydia website (http://www.chlamydialae.com)). There is little evidence to suggest differences in virulence among ocugential serovars, with a recent large study concluding that the severity of disease was not associated with a particular serovar or group of serovars. However, surveys of genital isolates from around the world have consistently shown that certain serovars occur more frequently than others, and that over time this pattern has remained constant. Specifically, in large surveys in the USA and Europe, serovars D and E account for approximately 50% of genital C trachomatis isolates, whereas serovar H is isolated from only 1–3% of culture confirmed infections. In addition, differences in the level of shedding among serovars have been noted in clinical specimens that roughly correlated with serovar prevalence.

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In women, 70–80% of infections are asymptomatic and repeated infection correlates with severe upper genital tract pathology, which in decreasing order of occurrence includes pelvic inflammatory disease, ectopic pregnancy, and tubal infertility. The factors that contribute to a particular outcome are unknown, but probably depend upon the interplay between the host’s complex response to infection and the equally complex, phenotypically variable, and environmentally responsive biphasic intracellular developmental cycle of chlamydia.

In attempts to understand the salient features of this interplay, a great deal of work has been carried out using slightly modified versions of a murine model of female genital tract infection first described by Tuffrey and Taylor-Robinson. Until recently, most of this work has been conducted using C muridarum, the mouse pneumonitis agent. However with the complete sequencing of C muridarum and several chlamydial strains representing human biovars of C trachomatis, studies are being conducted that compare the more clinically relevant phenotypes that might arise out of the variation seen at the genetic level. Noteworthy in the present context are reported differences between C muridarum and human ocugential serovars in sensitivity to interferon γ and elementary body (EB) associated cytotoxicity, with differences in interferon γ having a known effect on the course and outcome of female genital tract infection in mice. We used this model to determine whether the difference in prevalence among ocugential serovars might result from differences in virulence and the observed significant differences in the duration of infection among the various serovars that loosely corresponded to the clinical prevalence of the serovars, especially for the most and least prevalent serovars—D and E versus H and I. In addition, the invasiveness and level of bacterial shedding of the serovar D and H strains

Abbreviations: EB, elementary body; ifu, inclusion forming units; LDH, lactate dehydrogenase; MOI, multiplicity of infection; SP, sucrose phosphate transport medium
demonstrated that serovar D was able to establish culture confirmed infection of the uterine horns significantly more frequently, in addition to shedding more during the acute phase of infection. The factors that contribute to these distinct phenotypes are unknown.

The purpose of our study was to analyse the in vitro growth characteristics and EB associated cytotoxicity of the previously studied serovar D and H strains to identify differences that might explain the observed variation in the course of infection. Extrapolating the results of these studies could provide insight into the factors that influence serovar H prevalent and levels of shedding that are seen in human C trachomatis genital tract infection. Chlamydia muridarum was included in these analyses both to confirm the original observation relating to non-lipopolysaccharide EB cytotoxicity and to identify phenotypic traits that might explain the highly virulent and invasive nature of C muridarum infection in the mouse compared with the less virulent and invasive infection of human isolates in this model of human disease.

**MATERIAL AND METHODS**

**Chlamydia strains**

Polymerase chain reaction based restriction fragment length polymorphism serovar typed and mycoplasma free C trachomatis strains belonging to serovars D, H, and L2 and C muridarum was propagated and titrated in cycloheximide treated McCoy cell monolayers using standard techniques. EBs were purified using density gradient centrifugation, and concentrated stock suspensions in sucrose phosphate transport medium (SP) were frozen at −80°C.

**Growth cycle analysis**

Confluent McCoy cell monolayers (5 × 10⁶ cells) in 24 well tissue culture plates were inoculated with approximately 600 purified EBs. After centrifugation at 1500 ×g for one hour at 37°C, monolayers were washed and incubated at 37°C in fresh medium. Immediately after and at eight, 24, 32, 48, 56, and 72 hours post-infection, culture supernatants and monolayers suspended in SP were collected separately and in duplicate. Both types of specimens were sonicated and immediately frozen at −80°C until cultured for the presence and enumeration of inclusion forming units (ifu) in McCoy cell monolayers using iodine. A companion plate was used to verify the actual input inoculum, and to observe the kinetics and microscopic appearance of iodine staining inclusions throughout a 96 hour incubation period.

**EB associated cytotoxicity analysis**

EB associated cytotoxicity was assessed in confluent McCoy cell monolayers using two approaches: (1) visual evaluation of the cytopathic effect with light microscopy; and (2) measurement of changes in host cell membrane integrity using a lactate dehydrogenase (LDH) release assay (cytotoxicity detection kit; Roche, Indianapolis, Indiana, USA). As a control for possible cytopathic and membrane perturbing effects resulting from particulate burden and chlamydial lipopolysaccharide, a strain of C trachomatis L2 was included in our analysis. McCoy cells were grown overnight in 48 well tissue culture plates (2.2 × 10⁵ cells/well), and immediately before use the culture medium was replaced with serum free medium supplemented with 1% bovine serum albumin and containing rifampin (10 µg/ml) and doxycycline (5 µg/ml), to eliminate whole serum LDH activity and to prevent bacterial metabolism during inoculation. Purified EBs were diluted in SP and inoculated in duplicate on to McCoy cell monolayers at multiplicity of infection (MOI) ratios of 100 : 1, 25 : 1, 6.25 : 1, and 1.5 : 1. After centrifugation at 1500 ×g for one hour at 37°C, the supernatants were collected, and the monolayers were washed and incubated at 37°C in antibiotic containing serum free bovine serum albumin supplemented medium. Monolayers were inspected using standard light microscopy and subjectively scored for cytopathic effect immediately after inoculation, and when aliquots of culture medium were collected in duplicate at one, two, four, and six hours post-inoculation for the LDH assay, performed according to the manufacturer’s instructions.

**RESULTS**

**Growth cycle analysis**

Figure 1 shows the 72 hour replication and release kinetics assessed under standard culture conditions. During this period, C muridarum produced significantly more infectious units in the first 24 hours than were produced during the entire period for serovars D and H, and appeared to undergo two rounds of replication, as indicated by the decrease in the number of ifu isolated from monolayers between two peaks at 32 and 72 hours. In addition, C muridarum infected cells released ifu sooner—24 versus 48 hours—and in four to 10 times greater numbers than serovar D. In contrast, shedding of serovar H was first seen at 32 hours and, when adjusted for input, was both more sustained and equivalent to or greater in magnitude than the numbers of ifu released from C muridarum infected cells, resulting in the continued reduction in monolayer associated ifu and a coincidental increase in ifu recovered in the medium between 48 and 72 hours.

Microscopic observation of the iodine stained monolayers revealed C muridarum inclusions by 24 hours, uniformly increasing in size between 24 and 48 hours, followed thereafter by a decrease in the number of large inclusions and the appearance of an increasing number of small iodine staining inclusions, often in clusters. This resulted in a biphasic pattern of iodine staining inclusions (fig 1), with an initial maximum of 885 at 32 hours, which was used as the assigned value of the input inoculum, and a second peak of 860 at 72 hours, with both peaks corresponding to high points of ifu recovery from the infected monolayer (fig 1).

In contrast, inclusion development was similar for serovars D and H during the first 48 hours, with small iodine staining pronounced and release of Chlamydia trachomatis serovars D and H and C muridarum.
Differences between *Chlamydia trachomatis* and *C. muridarum*

**Figure 2** Kinetics of iodine staining inclusion development in McCoy cells.

Inclusions being first visible at 32 hours, followed by a relatively uniform increase in size up to 48 hours. However, beginning at 48 hours and continuing throughout the incubation period, serovar H infected monolayers contained inclusions that had apparently lost inclusion membrane integrity, and appeared as more diffuse, smudge-like inclusions within the cytoplasm of the cell.

The appearance of these distorted inclusions coincided with a decrease in the number of inclusions in the monolayer (fig 2) and an increase in ifu detected in the medium (fig 1). The maximum number of iodine staining inclusions for serovar D was seen at 56 hours (610) and for serovar H at 48 hours post inoculation (460). Between 48 and 96 hours of incubation, iodine staining inclusions for serovar H decreased to 41% of maximum at 56 hours to 8.6% at 96 hours; over the same period, serovar D dropped to 78% of maximum at 72 hours to 41% at 96 hours post-inoculation.

The net effect of these different replication and release phenotypes was the production of a significantly greater number of progeny/unit input for *C. muridarum* (950 : 1) compared with serovars D (235 : 1) and H (120 : 1).

**EB associated cytotoxin analysis**

Two methods were used to assess host cell cytotoxicity associated with purified EBs—direct microscopic assessment of the cytopathic effect and LDH release. Over time and in a dose dependent manner, McCoy cells inoculated with EBs of *C. muridarum* or either ocugentral serovar showed signs of toxicity, beginning with membrane perturbations at cell junctions, visible as early as one hour, and proceeding to varying levels of apparent lysis within two to four hours.

Although *C. muridarum* EBs were significantly more cytotoxic to McCoy cell monolayers at moderate (25 : 1) and high (100 : 1) MOI ratios compared with serovar D, serovar H EBs were significantly more cytotoxic than *C. muridarum* EBs, with an effect seen at the lowest MOI (1.5 : 1) tested, both microscopically (table 1) and in the more objective LDH release assay (table 2).

**DISCUSSION**

In an earlier study, we demonstrated a significant difference in virulence characteristics among strains representing seven ocugentral serovars of *C. trachomatis* in the murine model of female genital tract infection. In a comparison of the most (serovar D) and least (serovar H) virulent strains in this collection, the serovar D strain established a longer and more invasive infection, which was characterised by more shedding during the acute phase of infection. These findings suggested that the differences in the prevalence and shedding among serovars observed in humans might be related to differences in the ability to establish more or less durable infection, which may contribute to differences in genital tract invasiveness.

In our in vitro study, we assessed the growth characteristics and EB associated cytotoxicity of the serovar D and H strains previously characterised in the mouse, and showed that serovar H EBs were cytotoxic at an MOI of approximately 1 EB/McCoy cell, whereas a comparable level of cytotoxicity was seen at an MOI of 25 for the serovar D strain. This phenotypic difference, together with the possibly linked differences in replication and release kinetics, might help explain the differences in both the course of infection between these strains in the mouse, and the incidence and level of shedding between serovars D and H in human genital tract infections.

Based on the expression data presented in the original description, it can be hypothesised that the cytotoxin plays a role throughout the course of infection. It is possible that the more cytotoxic serovar H EBs might cause the death of some host cells upon contact, explaining the larger infectious dose required to establish infection in mice. During the replication cycle, newly produced cytotoxin could alter the integrity of both the inclusion and cytoplasmic membranes, causing earlier disruption of the replication cycle and premature lysis of infected cells. This would explain both the earlier and...
larger number of ifu released into the medium, the appearance of smudge cells, the disappearance of more cells with intact inclusions from the monolayer, and the production of less total ifu during the course of in vitro infection with serovar H. During genital tract infection, this would result in the release of less ifu, as previously described in both mice and humans. Ultimately, less shedding would reduce the number of ifu available to sustain infection, which might contribute to a shorter and less invasive infection, as seen in the mouse model, and a reduced level of transmission among humans, which would explain the well documented difference in the prevalence of infection with these serovars in human populations from around the world. The ability to produce large numbers of infectious units rapidly probably contributes to the ability of C muridarum to ascend the genital tract and ultimately deliver infectious challenge to the upper genital tract within the first seven days of infection, which results in the severe pathology seen during C muridarum infection. One can speculate that in the process C muridarum is capable of avoiding the innate immune system during each round of replication, being ultimately confined and contained by the aggressive inflammatory and immune response that occurs either just before or coincidental with the involvement of the entire genital tract, which in this last instance results in infertility.

Although compelling in its focus on the most severe sequelae associated with infection, C muridarum infection of the mouse female genital tract is not, in many of its features, analogous to human infection with oculogenital serovars of C trachomatis, particularly in the nature of the severe upper genital tract pathology it is meant to mimic. This infrequent and previously overestimated outcome associated with infection is thought to be a chronic process associated with either persistent or multiple infections, in which host

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<th>Assessment of LDH release from McCoy cell monolayers after inoculation with chlamydia EBs</th>
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<td>Time after inoculation (hours)</td>
<td>% LDH release from McCoy cell monolayers after inoculation with chlamydia EBs at the indicated MOI*</td>
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*Confluent McCoy cell monolayers were infected at MOIs of 1.5, 6.25, 25, and 100, and incubated at 37°C in the presence of 10 μg/ml of rifampicin and 5 μg/ml of doxycycline. LDH release was assayed in the supernatant immediately after inoculation and in aliquots taken at 1, 2, 4, and 6 hours post-inoculation. The percentage release was calculated using the total cell LDH activity released during the complete lysis of uninoculated monolayers as 100%. Spontaneous release from uninoculated monolayers was negligible at all time points.

EB, elementary body; LDH, lactate dehydrogenase; MOI, multiplicity of infection.

Take home messages

- Differences in elementary body associated cytotoxicity between Chlamydia trachomatis serovars D and H are linked with in vitro growth characteristics that can explain both the shorter and less invasive course of serovar H infection in the mouse female genital tract, and more importantly the higher prevalence and greater level of shedding that have been seen in human infections with serovar D.
- Differences seen between C muridarum and C trachomatis serovars D and H, particularly in the much faster in vitro replication and release kinetics, correspond to the more rapid and invasive course of mouse genital tract infection with C muridarum compared with human oculogenital serovars.
- Taken together with previously published reports, these findings provide an objective basis to question the appropriateness of C muridarum as a model strain in the murine model of human female genital tract infection, and establish the model’s capacity to assess the potential effects of phenotypic differences among human isolates on the course and outcome of human infection.
susceptibility factors play a role, and not the result of inflammatory processes associated with a single acute disease episode. In contrast, infection of the mouse with human strains mimics in many ways both the course and outcome of infection in most women—that is, an asymptomatic and self limiting infection that only rarely results in severe upper genital tract sequelae. The results presented in our study, together with other reports describing differences in potentially virulence determining phenotypes between C. muridarum and human isolates, again raise the issue of the appropriateness of C. muridarum as a model agent in the murine model of female genital tract infection. The issue is made more compelling given that the collection of phenotypes that define C. muridarum may not have a single counterpart in the diverse collection of serovars that comprise the human ocuogential biovar of C. trachomatis. Systematic in vitro and in vivo investigation of the phenotypic variation among human disease causing strains could provide results with translational value that will help direct the development of a vaccine and/or other intervention strategies effective against C. trachomatis genital tract infections.

In conclusion, the results of our study provide a strain specific phenotype based explanation for the profound differences in the course and outcome of female genital tract infection in mice infected with strains belonging to the human ocuogential biovar of C. trachomatis and C. muridarum, raising the issue of the appropriateness of using C. muridarum as a surrogate for C. trachomatis in this model of human disease. More importantly, the human epidemiological data relating to differences in the prevalence and level of shedding that occurs during infection with ocuogential serovars D and H can be understood in the context of defined phenotypic characteristics.

References