Chlamydiophila psittaci in homing and feral pigeons and zoonotic transmission

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Chlamydiophila psittaci is a zoonotic disease in birds caused by Chlamydiophila psittaci, an obligate intracellular bacterium. There are seven known avian outer-membrane protein A (ompA) genotypes designated A–F and E/B. The importance of genotyping lies in the fact that certain genotypes tend to be associated with certain hosts and a difference in virulence. Genotype B is the most prevalent in pigeons, but the more virulent genotypes A and D have also been discovered. The current study assessed the prevalence of C. psittaci in 32 Belgian homing-pigeon facilities and in 61 feral pigeons captured in the city of Ghent, Belgium. Additionally, zoonotic transmission of C. psittaci was investigated in the homing-pigeon facilities. Homing pigeons were often infected, as at least one of the lofts was positive in 13 of the 32 (40.6%) pigeon breeding facilities. Genotypes B, C and D were detected. Zoonotic transmission was discovered in 4 of the 32 (12.5%) pigeon fanciers, revealing genotype D in two of them, whilst genotyping was unsuccessful for the other two human pharyngeal swabs. This study clearly demonstrates the possible risk of C. psittaci zoonotic transmission from homing pigeons. Pigeon fanciers often (37.5%) used antibiotics for prevention of respiratory disease. Because of the risk of developing drug-resistant strains, regular use of antimicrobial drugs must be avoided. This study is believed to be the first to detect C. psittaci in Belgian feral pigeons. The prevalence rate in the city of Ghent was extremely low, which is beneficial for public health.

INTRODUCTION

Chlamydiophila psittaci is a zoonotic disease in birds caused by Chlamydiophila psittaci, an obligate intracellular bacterium. Symptoms include abnormal respiratory signs, nasal discharge, diarrhoea, polyuria and dullness (Vanrompay et al., 1995a). Zoonotic transmission happens through direct contact or inhalation of infected aerosols. The clinical course in humans varies from asymptomatic to severe systemic disease with interstitial pneumonia. Symptoms frequently include high fever (up to 40.5°C) accompanied by a relatively low pulse, chills, headache, myalgia, non-productive coughing and difficulty breathing (Beeckman & Vanrompay, 2009). The incubation period ranges from 5 to 14 days.

Currently, there are seven C. psittaci outer-membrane protein A (ompA) genotypes designated A–F and E/B. The importance of genotyping lies in the fact that certain genotypes tend to be associated with certain hosts and differ in virulence. Genotypes A and D are more virulent than genotype B (Vanrompay et al., 1994, 1995b). Genotypes A–E and E/B have been found in pigeons (Geens et al., 2005; Laroucau et al., 2008; Sayada et al., 1995). However, in Belgium and neighbouring countries, only genotypes A, B and D have been discovered so far (Vanrompay et al., 1993, 1997), with genotype B being the most prevalent (Magnino et al., 2009). Recently, genotype B has been discovered in three human samples in The Netherlands and may be an underestimated source of disease (Hedema et al., 2006b).

In 1940, C. psittaci was isolated for the first time from homing pigeons (Pinkerton & Swank, 1940). Since then, several serological surveys have been conducted in pigeons. However, most articles have focused on C. psittaci in feral pigeons (Columba livia). Feral pigeons are more commonly known as ‘urban’ or ‘city’ pigeons. Thirty-eight studies on the seroprevalence of C. psittaci in feral pigeons conducted from 1966 to 2005 revealed a seropositivity rate ranging from 12.5 to 95.6% (Haag-Wackernagel, 2005; Laroucau...
et al., 2005; Mitevski et al., 2005; Prukner-Radovčić et al., 2005; Tanaka et al., 2005). More recent serological studies performed in feral pigeons in Italy, and Bosnia and Herzegovina, showed a seropositivity of 48.5 and 26.5%, respectively (Ceglie et al., 2007; Ilieski et al., 2007). Fourteen studies on the culture of C. psittaci conducted in European feral pigeons from 1980 to 2007 revealed 1.6 to 57% positivity (Magnino et al., 2009). The highest percentage of culture positives was found in Paris (Trap et al., 2009). Eleven studies using nucleic acid amplification assays performed from 2003 to 2007 demonstrated the presence of chlamydial DNA in 3.4–52.6% of the feral pigeons examined (Magnino et al., 2009; Vázquez et al., 2010). Data on the occurrence of C. psittaci from 35.9 to 50.1% (Alexander et al., 1992) for culture and 2 ml DNA stabilization buffer for nested PCR and one in chlamydial transport medium (Vanrompay et al., 1992). Incompletely returned packages were excluded from the study.

In the same period, a study of 61 feral pigeons was conducted taking two pharyngeal swabs from each individual pigeon. The pigeons were captured on several days during the winter period of 2008 in the city of Ghent, Belgium. The captured pigeons were all adults and of both sexes. After capture, the pigeons were euthanized as part of pest control. The swabs were taken from the euthanized animals. Again, one swab was placed in DNA stabilization buffer for nested PCR and one in chlamydial transport medium for culture. All swabs were kept on ice during transport and stored at –80°C until use.

C. psittaci nested PCR. Bacterial DNA was prepared by the standard extraction method, as described previously (Van Loock et al., 2005b). The DNA was purified further by extraction in phenol/chloroform/isoamylalcohol (25:24:1, v/v/v; Invitrogen). Purified DNA was precipitated by adding ethanol (100%) and sodium acetate (5 M), and incubated at –80°C for 45 min. The DNA was centrifuged for 15 min (15490 g) at 4°C. Pellets were washed with 70% ethanol and suspended in 50 μl MilliQ water.

The presence of the C. psittaci ompA gene was examined using a C. psittaci-specific nested PCR and the results analysed by agarose gel electrophoresis (Van Loock et al., 2005b). Briefly, nested PCR generated a product of 472 bp. Negative samples were tested to exclude false negatives by adding 10 ng internal inhibition control to assess possible polymerase inhibition. The inhibition control consisted of a pcDNA1 vector in which the ompA gene of C. psittaci, inserted as a 231 bp fragment, was cloned (Van Loock et al., 2005b).

ArrayTube DNA microarray for ompA genotyping. For C. psittaci genotype determination, sample DNA was amplified and biotin-labelled ompA fragments of 418 and 530 bp were generated for the ArrayTube microarray (Clondiag Chip Technologies) using 40 cycles of 95°C for 30 s, 50°C for 30 s and 72°C for 30 s, with primers VD1-F (5′-ACTACGGAGAATTGTTTTCGATCGTGT-3′), Bio-VD2-r (5′-biotin-GTGGCAGACATATGTTTTCGATCGTGT-3′), Bio-VD2-r (5′-GGGCWGGMTTCCATAYGCICARTC-3′) and Bio-ompA-rev (5′-biotin-TCCCTTAGAATCTGAATTGAGC-3′). The results were visualized by agarose gel electrophoresis (Sachse et al., 2008). Hybridization was performed at 53°C for 60 min. Subsequently, the ArrayTube was incubated with streptavidin-conjugated horseradish peroxidase (ArrayTube staining kit; Clondiag). Finally, peroxidase substrate (3,3′,5,5′-tetramethyl benzidine derivative from the ArrayTube staining kit) was added. Hybridization signals were analysed using an ATR-03 array tube reader (Clondiag) and Iconoclust software version 2.3 (Clondiag).

C. psittaci culture. Samples that were positive by PCR or showed signs of inhibition were tested by culture. The presence of viable C. psittaci was examined by isolation in buffalo green monkey kidney cells identifying the organism by a direct immunofluorescence staining (IMAGEN; Oxoid) at 6 days post-inoculation, as described previously (Vanrompay et al., 1992). The number of C. psittaci-positive cells was counted (magnification ×600; Nikon Eclipse TE2000-E microscope) and a score from 0 to 4 was given (Table 1).
Table 1. Culture scoring system

<table>
<thead>
<tr>
<th>Score</th>
<th>Meaning</th>
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<tbody>
<tr>
<td>0</td>
<td>No EBs or inclusions</td>
</tr>
<tr>
<td>0.5</td>
<td>1–3 EBs</td>
</tr>
<tr>
<td>1</td>
<td>&gt;3 EBs and/or 1–2 IPCs</td>
</tr>
<tr>
<td>1.5</td>
<td>3–5 IPCs</td>
</tr>
<tr>
<td>2</td>
<td>&gt;5 IPCs in less than a quarter of the fields</td>
</tr>
<tr>
<td>2.5</td>
<td>IPCs in a quarter of the fields</td>
</tr>
<tr>
<td>3</td>
<td>IPCs (small inclusions) in half of the fields</td>
</tr>
<tr>
<td>3.5</td>
<td>Large IPCs in half of the fields</td>
</tr>
<tr>
<td>4</td>
<td>IPCs in all fields</td>
</tr>
</tbody>
</table>

EB, Elementary body; IPC, inclusion-positive cell.

Statistics. The data were examined statistically using a non-parametric test, as the sample populations were smaller than 30. The medical questionnaire enquired about respiratory complaints in the human participants, such as wheezing, dyspnoea, shortness of breath, sore throat, chest pain, runny nose, coughing, sneezing and coughing up mucus. For each of these complaints, the reporting frequency was scored as follows: 0, when the complaint was never reported; 1, if the complaint was reported once; 2, if it was reported several times; and 3, if the complaint was regularly reported. Scores were added to attain one overall respiratory complaint score per person. The same scoring method was applied to the reported respiratory symptoms and conjunctivitis in pigeons. The questionnaire also enquired about treatment with antibiotics in the pigeons as well as in the pigeon fanciers. A score of 0 or 1 was given when no treatment or recent treatment (in the last 3 months) was given, respectively.

RESULTS AND DISCUSSION

**C. psittaci** in homing pigeons

A total of 32 (61.5 %) out of 52 breeding facilities returned complete packages, resulting in samples from 96 lofts (32 young, 32 breeder and 32 racing-pigeon lofts) and 32 humans. The remaining 20 facilities returned an incomplete package and were therefore excluded from further examination. The results are presented in Table 2. A total of 2 of the 32 lofts (6.3 %) of young pigeons were PCR positive, whilst 11 of the 32 samples (34.4 %) contained PCR inhibitors. None of the 32 samples of the breeder pigeons were positive, whilst 9 (28.1 %) contained polymerase inhibitors. Adult racing pigeons were all PCR negative. However, 10 of the 32 samples (31.3 %) contained polymerase inhibitors and could not be examined by PCR. Overall, PCR inhibition was frequently observed when examining floor swabs, as 30 of the 96 floor samples (31.3 %) contained enzyme inhibitors.

All positive samples and those with inhibitors (a total of 32 samples) were examined by culture. Nine out of thirteen (69.2 %), two out of nine (25 %) and seven out of ten (70 %) of the young, breeder and racing-pigeon lofts, respectively, were culture positive. All PCR-positive samples contained viable *C. psittaci*. The mean isolation scores for young, breeder and racing-pigeon lofts were 2.17, 1.25 and 1.5, respectively. Lofts were considered positive if PCR or culture was positive. Nine (28.1 %), two (6.3 %) and seven (21.9 %) of the thirty-two young, breeder and racing-pigeon lofts were positive, respectively.

Lofts of young birds were more often positive, showing higher mean culture scores, followed by lofts of racing and finally breeder pigeons. Young and racing-pigeon lofts were found to be positive significantly more often than breeder-pigeon lofts, but the mean culture scores were not significantly different. However, mean culture scores need to be interpreted with care, as the amount of faeces and medium present in the returned swab tubes differed. This inconvenience is inherent to the methodology of the do-it-yourself sampling package that was used. Nevertheless, our results for young birds are in accordance with other studies demonstrating more frequent *C. psittaci* infections in young birds (Bedson, 1955; Schettler *et al.*, 2001). Young birds are partially protected by maternal antibodies. As soon as maternal antibody titres decline, young birds may experience a primary infection, which is often accompanied by clinical signs due to the absence of an immunological memory (Van Loock *et al.*, 2005a).

Racing-pigeon lofts were more often positive than breeder-pigeon lofts, showing higher mean culture scores. Racing pigeons are transported, which creates stress, and crowding augments the infection pressure. During races, contact with wild birds, a reservoir for *C. psittaci*, may occur. Tiring could also render the homing pigeons more prone to infection. Whilst breeder pigeons suffer reproductive stress, which could render them more prone to infection, our results indicated that young and racing pigeons are more sensitive to *C. psittaci* infection (Andersen & Vanrompay, 2003). Overall, in 13 (40.6 %) of the 32 pigeon breeding facilities, at least one of the floor swabs was positive.

Table 2. *C. psittaci* detection in pharyngeal swabs of pigeon fanciers and in floor swabs of pigeon lofts

<table>
<thead>
<tr>
<th>Test</th>
<th>No./total (%)</th>
<th>Young pigeons</th>
<th>Breeder pigeons</th>
<th>Racing pigeons</th>
<th>Humans</th>
</tr>
</thead>
<tbody>
<tr>
<td>PCR positive</td>
<td>2/32 (6.3)</td>
<td>0/32 (0)</td>
<td>0/32 (0)</td>
<td>1/32 (3.1)</td>
<td></td>
</tr>
<tr>
<td>Inhibition in PCR</td>
<td>11/32 (34.4)</td>
<td>9/32 (28.1)</td>
<td>10/32 (31.3)</td>
<td>5/32 (15.6)</td>
<td></td>
</tr>
<tr>
<td>Culture positive</td>
<td>9/13 (69.2)</td>
<td>2/9 (25)</td>
<td>7/10 (70)</td>
<td>4/6 (66.7)</td>
<td></td>
</tr>
</tbody>
</table>
demonstrating the high prevalence of *C. psittaci* in Belgian homing pigeons.

Thirteen out of eighteen positive floor samples (72.2 %) could not be genotyped directly on clinical samples. They all contained polymerase inhibitors generating negative microarray results. They were positive by culture but unfortunately did not grow efficiently enough to extract sufficient DNA for the genotyping microarray. Samples of five young-pigeon lofts were successfully genotyped, revealing *ompA* genotypes B (three lofts), C (one loft) and D (one loft). So far, genotype C has been detected only in a Japanese pigeon (Sayada *et al.*, 1995). This genotype is found mostly in waterfowl. It is highly pathogenic to domestic waterfowl, and is an admitted occupational hazard for slaughterhouse workers and generally for all people in contact with ducks.

A *C. psittaci*-positive result in birds was not statistically correlated with the presence or absence of respiratory disease (*p*=0.821), nor with antibiotic administration (*p*=0.563). *C. psittaci* can be detected in carriers showing no clinical disease. In addition, respiratory disease can also be due to other systemic bacterial infections, viral infections such as herpesviruses and even fungal infections (Marlier & Vindevogel, 2006).

As in psittacine breeding facilities, antibiotics were also used prophylactically (Harkinezhad, 2008). In our study, 12 of the 32 pigeon fanciers (37.5 %) used tetracyclines for prevention of respiratory disease. In Belgium, a prescription by a veterinarian is required to obtain these drugs. However, tetracyclines are sold on the Internet without a prescription as a prescription is not needed in every country. A vaccine as well as information on the sensible use of antimicrobial drugs in homing pigeons are needed to prevent psittacosis in humans and the development of drug-resistant chlamydial strains.

### *C. psittaci* zoonotic transmission from homing pigeons to humans

Only 1 of the 32 pigeon fanciers (3.1 %) tested positive by nested PCR. A total of 5 of the 32 human pharyngeal samples (15.6 %) contained polymerase inhibitors. The one PCR-positive sample and the five containing inhibitors were examined for the presence of viable *C. psittaci*. Four of the six samples (66.7 %) were positive by culture, including the one that was positive by nested PCR, with a mean isolation score of 2.13. Overall, 4 of the 32 pigeon fanciers (12.5 %) were positive for *C. psittaci*. Two of the four positive samples (50 %) could be genotyped. They both revealed genotype D, which is a virulent genotype often found in turkeys and known to be transmitted to humans (Van Droogenbroeck *et al.*, 2009). One person testing positive had been treated with doxycycline until 2 weeks before the start of the study. He suffered from respiratory disease (sneezing, coughing up mucus and head cold). His pigeons had respiratory symptoms and were still being treated with doxycycline. Probably due to this treatment, neither viable *C. psittaci* nor chlamydial DNA could be detected in the loft samples. This pigeon fancier had no contact with other birds. The other person testing positive reported mild respiratory symptoms (shortness of breath and sneezing) but took no medication. Although his pigeons showed respiratory problems and were being treated with enrofloxacin, we detected chlamydial DNA as well as viable *C. psittaci* genotype D. This pigeon fancier was also infected with genotype D.

Harkinezhad *et al.* (2009b) showed that 7 % of apparently healthy people having daily contact with homing pigeons were positive by PCR and/or by a recombinant major outer-membrane protein antibody ELISA. Our study revealed a slightly higher percentage (12.5 %) of positive persons. The relatively small number of humans included in both studies (both ~30) makes a robust comparison difficult. However, both studies indicated the risk of zoonotic transmission from homing pigeons to humans.

Interestingly, infection in pigeon fanciers occurred statistically more frequently when pigeons experienced respiratory disease and were not simply *C. psittaci* positive (*p*=0.042). Thus, the presence of positive animals in itself did not lead to a higher infection rate in humans (*p*=0.648). Respiratory disease in pigeons most often reflects an acute infection accompanied by excretion of a higher amount of infectious *C. psittaci* than in birds without respiratory symptoms, which most often have intermittent shedding (Harkinezhad *et al.*, 2009a).

Three of the four culture-positive persons (75 %) showed respiratory symptoms. However, 10 of the 28 PCR- and culture-negative persons (35.7 %) also showed respiratory symptoms. Respiratory symptoms due to *C. psittaci* can easily be confused with pigeon fanciers' lung, which is due to inhalation of organic dust material (Calvert *et al.*, 1999).

### *C. psittaci* in feral pigeons

One of the sixty-one feral pigeons (1.6 %) from the city of Ghent was PCR positive and three of the samples (4.9 %) contained polymerase inhibitors. The PCR-positive sample and those containing inhibitors were subsequently examined by culture. Only the PCR-positive sample contained viable *C. psittaci* with a low culture score of 2. Passages on buffalo green monkey kidney cells and subsequent genotyping were unsuccessful. The latter was probably due to an insufficient amount of chlamydial DNA, rather than to the presence of a new *ompA* genotype.

A study in Amsterdam pigeons, conducted in the low breeding period, revealed 5 % PCR-positive pigeons (Hedema *et al.*, 2006a). Our study, also conducted when breeding activity was low, revealed only one *C. psittaci*-positive pigeon, which is comparable to the low infection rate in the city of Amsterdam. This low infection rate is beneficial to public health. However, caution is needed, as zoonotic transmission of *C. psittaci* from feral pigeons is
known to be an underestimated source of infection (Haag-Wackernagel & Moch, 2004). Other studies on feral pigeons in the city of Ghent should be conducted, including at different time periods (e.g. the high breeding season) and other locations (with high concentrations of pigeons). Feral pigeon populations should be managed carefully in the urban environment to obtain an appropriate-sized and healthy population.

For this study, pharyngeal swabs were used instead of faecal or cloacal swabs. Whilst this makes comparison with the pigeon breeder study less easy, this technique does not have the disadvantage of intermittent shedding (Harkinezhad et al., 2009a). When cloacal swabs are taken, false negatives can be obtained because of intermittent shedding. We were unable to genotype the PCR-positive sample. In the city of Amsterdam, ten genotyped PCR-positive samples were all genotype B. Genotype B may be an underestimated source of zoonotic infections and human disease (Hedema et al., 2006b).

Conclusions

We found that homing pigeons were often infected, as at least one of the lofts was positive in 13 of the 32 pigeon breeding facilities (40.6%). Genotypes B, C, and D were detected. Zoonotic transmission was discovered in 4 of the 32 pigeon fanciers (12.5%), revealing genotype D in 2 of them, whilst genotyping was unsuccessful for the other 2 human pharyngeal swabs. This study demonstrates the possible risk of C. psittaci zoonotic transmission from homing pigeons. Further research on more cases is necessary to confirm this result. The pigeon fanciers often (37.5%) used antibiotics for prevention of respiratory disease. Because of the risk of developing drug-resistant strains, as described for Chlamydia suis (Dugan et al., 2004), regular use of antimicrobial drugs must be avoided. Our study is believed to be the first to detect C. psittaci in Belgian feral pigeons. However, the prevalence rate in the city of Ghent was extremely low, which is beneficial for public health.

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Epidemiological investigation of mapping for avian Chlamydia psittaci


