Declining Mortality in American Crow (Corvus brachyrhynchos) Following Natural West Nile Virus Infection

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SUMMARY. The American crow (Corvus brachyrhynchos) is known to suffer 100% mortality from infection with the New York 1999 strain of West Nile virus (WNV). Following the initial detection of WNV in North America in 1999, we measured prevalence of WNV-reactive antibodies ("seroprevalence") in free-ranging American and fish crows (Corvus ossifragus) of central New Jersey after each transmission season through 2005. In 2002, seroprevalence in American crow juveniles increased to 14% from the 5% of the previous year, potentially indicating increased survival in this species. Using the annual seroprevalence measurements and the number of human West Nile neuroinvasive disease cases as a surrogate for WNV transmission intensity, we developed a model to estimate the annual WNV-associated mortality rates among both of these crow species. Our model supports the hypothesis that mortality is changing over time; the WNV-associated mortality rate declined over time by 1.5% for American crow and by 1.1% for fish crow. The probability that the trend in mortality was negative was 90% for the American crow and 60% for the fish crow.

Key words: West Nile virus, crows, corvids, seroprevalence, infection, mortality

Abbreviations: PRNT = plaque reduction neutralization test; WNND = West Nile neuroinvasive disease; WNV = West Nile virus

West Nile virus (WNV), a pathogenic arbovirus with avian primary hosts and culicid vectors, is associated with significant corvid mortality in North America (25). Numerous corvids are susceptible to WNV, including the American crow (Corvus brachyrhynchos) in particular, but also fish crow (Corvus ossifragus), two species of magpies (Pica spp.), and several species of jays (e.g., Cyanocitta spp.) (10,19,20,32). Prior to the introduction of WNV in the Western Hemisphere in 1999, large numbers of crow deaths due to infectious diseases were unusual. Since 1999, dead crows have tested positive for acute WNV infection at alarming rates (2,27), and clusters of dead crows have spatially predicted human cases of WNV-induced disease (15,30,32).

Corvid mortality quickly became a hallmark of the North American WNV invasion and was promoted as an early warning signal for local WNV activity (14,16). Experimental infections of American crows with the New York 1999 strain revealed that 100% of infected crows succumbed and that crows became infected from each other in the absence of mosquito vectors (20,26), an observation that has been corroborated in the field (13). The New York 1999 strain was found to contain a bird-virulent mutation that permitted replication at the high body temperatures reached by febrile crows and other birds (4,5). Local WNV transmission to American crows was so intense in the eastern United States (3) that some local populations declined by almost 70% (9,35).

In spite of the substantial die-offs of crows associated with WNV, avian mortality surveillance seemed to gradually lose sensitivity for early detection of WNV, relative to other surveillance activities (6,28). The sensitivity of avian mortality surveillance for detecting WNV may decrease over time due to diminishing populations of WNV-sensitive bird species, to an accumulation of surviving birds that resist infection...
due to innate and acquired immunity and, in the case of species with high mortality rates, to selection of mortality-resistant genotypes (18). However, these explanations have not been proven in the case of the WNV epizootic in North America. We sought to corroborate the decrease in sensitivity of corvid-based avian mortality surveillance for WNV by evaluating the survival rates among crows naturally infected with WNV in New Jersey in the years following WNV introduction. Accordingly, we determined WNV-reactive antibody prevalence ("seroprevalence") rates among free-ranging American and fish crows following each WNV transmission season from 1999 through 2005 in central New Jersey. We then used West Nile neuroinvasive disease (WNND) incidence in humans as a measure of transmission intensity and, therefore, as an indicator of crow infection rates that would be independent of crow mortality. Given observed seroprevalence rates and this human case-derived infection rate, we could then estimate annual mortality rates to evaluate concurrent changes in mortality (and thus, survival) over time.

**MATERIALS AND METHODS**

American and fish crows were trapped at the solid waste facilities at Middlesex County Utilities Authority (Edgeboro), East Brunswick, New Jersey. This 300-ac landfill, located near the Raritan Bay, receives about 2600 solid-trash tons/day and is located about 52 km SW from the Borough of Queens, New York, where human WNND cases were first reported in the United States. The trap was located close to compost piles adjacent to the landfill. A previous study demonstrated that crows from at least 4 km away visited the compost piles (8). Trapping took place from late fall (beginning in late October) through early spring (ending in early April) of each year. Although traps were in place throughout the fall, winter, and spring of each year, the number of times crows were successfully caught varied considerably. This resulted in a wide variation of birds being caught and, subsequently, the latter years are represented by fewer birds and, thus, more uncertainty for the seroprevalence measures.

Crows were caught using a drop-in style trap, commonly called a Scandinavian or Australian crow trap (7). Crows were identified to species and age based on tarsus or culmen length (31), voice, or P9/P5 ratios among the primary feathers of the wing (24) and then banded with a United States Geological Survey band. A 2-ml sample of blood was taken via jugular venipuncture, allowed to coagulate for 30 min, and placed on ice. After sampling, the birds were held for 20 min to ensure that the puncture areas were not bleeding and then released nearby. Samples were centrifuged at 1288 × g at 4 C for 30 min, to separate red blood cells from serum, and stored at −70 C. Prior to 2003, samples were analyzed by a plaque reduction neutralization test (PRNT) for antibody detection (1). Beginning in 2003, samples were screened for antibodies using a WNV-specific blocking enzyme-linked immunosorbent assay (17). Positive samples were confirmed using PRNT. Samples were collected according to established Biohazard Safety–Animal protocols at Rutgers University.

To test the hypothesis that mortality is changing over time, we adopted a Bayesian approach to analyze the relationship between seroprevalence in juvenile birds, , after transmission season, , as a function of WNV incidence, , and WNV-associated mortality, , as described by Komar et al. (21):

\[ i_t = \frac{i_t - i_t m_t}{1 - i_t m_t} \]

The Bayesian approach allowed us to estimate these parameters while considering the uncertainty in each of them. They were estimated by simulation in WinBUGS (Version 1.4; MRC Biostatistics Unit, Cambridge, U. K. and Imperial College School of Medicine, London, U. K.) (23) (code is available from the authors upon request). The seroprevalence, , is estimated from a binomial distribution,

\[ j_t \sim \text{Binomial}(s_t, n_t) \]

where , is the number of seropositive juvenile birds out of those tested, , Juvenile birds are used exclusively in the model for every transmission season after 1999, because they are assumed to be universally susceptible to infection at the beginning of the transmission season. All birds were used for 1999 because the virus was not previously present in the United States and, therefore, no adult birds would have had pre-existing immunity.

Initial mortality, , was estimated for each species based on data from laboratory-infected crows (5,20,26,33),

\[ d \sim \text{Binomial}(m_t, n) \]

where is the number of birds that died out of the number infected, . Because our aim was to detect a trend in mortality over time, we modeled mortality as a linear function of time,

\[ m_t = m_0 + \Delta_m(t - 1), \]

where is the yearly change in mortality.

We used the reported number of WNND cases in humans, , as an indicator of corvid infection rate, . The incidence of human neuroinvasive disease is modeled as a Poisson distribution,

\[ c_t \sim \text{Poisson}(\lambda_t), \]

where avian incidence is assumed to be a function of its mean, ,

\[ \lambda_t = \alpha x, \]

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where is a non-zero, positive real number. New Jersey human WNND cases were obtained from Arbonet (Centers for Disease Control and Prevention, unpublished data).

**RESULTS AND DISCUSSION**

WNV-specific antibodies were detected annually (1999–2005) in American and fish crows in central New Jersey following the introduction of WNV into the United States (Table 1). Seroprevalence was generally higher for fish crow versus American crow, a pattern also observed in the southern United States (34) and likely explained by a higher WNV-associated mortality rate in American crows. Experimental infection of mature crows with the New York 1999 strain of WNV resulted in approximately 55% mortality for fish crow and complete mortality for American crow (5,20,26,33). The initial low seroprevalence (<=4%) in 1999 and 2000 may have resulted from low population exposure to virus, high mortality rates, or both. Indeed, large numbers of crow carcasses of both species were observed in New Jersey during this period, and many of them tested positive for WNV infection (29). Seroprevalence was higher for juveniles of both crow species in 2002, suggesting greater transmission intensity and an increase in survival rate for the American crow. A slight reduction in the virulence of the virus, or an evolutionary selection for genetically adapted crows, could result in increasing antibody seroprevalence rates over time, as more individuals survive infection.

Because the WNV-induced mortality rate of crows in nature is unknown and possibly changing, we used a simple mathematical model that links three variables: seroprevalence, infection rate, and mortality. Mortality is calculated as a function of seroprevalence and infection rate. Because only seroprevalence was directly measured, we used a surrogate for infection rate: the number of human WNND cases. WNND incidence provides a good measure of the intensity of WNV transmission for this region across all years of this study because WNND cases are assumed to be universally susceptible to WNV infection at the beginning of the transmission season. All birds were used for 1999 because the virus was not previously present in the United States and, therefore, no adult birds would have had pre-existing immunity. Initial mortality, , was estimated for each species based on data from laboratory-infected crows (5,20,26,33),

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increased probability of a mortality reduction is illustrated in Figure 2. The annual median decrease in mortality was approximately 1.5% per year (95% credible interval, [-44x-44]0.6%, 7.6%) and 1.1% per year [54x54]5.4%, 9.7%] for American and fish crow, respectively. Decreased mortality could be due to increased resistance in the crow population, or to attenuation of WNV itself. Both hypotheses warrant further investigation. Interestingly, a viral-strain shift was observed in the northeast United States in 2002 (12). Although the effects of this strain shift on crow pathogenesis, immune response, and survival are unknown, this 2002 strain was observed to be slightly attenuated in mice (11).

Our study serves to demonstrate the relationship between seroprevalence and mortality rate. These two values vary independently, depending on separate biologic processes. As an example, juvenile fish crow seroprevalence increased after the high transmission seasons of 2002 and 2003 and dropped again in 2004. The pattern of changing seroprevalence in fish crow likely reflects the general changes in WNV activity, rather than a significant change in mortality rate. Other factors that may contribute to variation in avian seroprevalence include attrition of local birds (due to WNV), movement of birds from areas with different WNV activity, or mutations in the WNV genome leading to variations in transmissibility.

The number of carcasses of both species sent in by the public dropped over the study period (29). Sources of this decrease may include decreased public participation, a smaller corvid population and, as our paper reports, an increase in American crow survival. LaDeau et al. (22) found that the introduction of WNV significantly reduced American crow, but not fish crow, abundance through 2005, suggesting that the smaller crow population may be the primary factor leading to fewer crow-carcass submissions. Although declining mortality in crows signals a likely reduction of sensitivity of avian mortality surveillance for WNV, the decline we detected was on the order of 10% for American crows.

Table 1. West Nile virus antibody prevalence rates (%) in American and fish crows, by age, in New Jersey 1999–2005. Sample size (n) and confidence intervals (95% CI) are given below each seroprevalence.

<table>
<thead>
<tr>
<th>Winter following year</th>
<th>American crow</th>
<th>Fish crow</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Adult % (n/95% CI)</td>
<td>Juvenile/naïve % (n/95% CI)</td>
</tr>
<tr>
<td>1999</td>
<td>2.3A (43/0.4–12.1)</td>
<td>3.7A (27/0.7–18.3)</td>
</tr>
<tr>
<td>2000</td>
<td>2.5 (40/0.4–12.9)</td>
<td>0 (7/0–35.4)</td>
</tr>
<tr>
<td>2001</td>
<td>14.3 (14/4.0–39.9)</td>
<td>14.3 (8/2.2–47.1)</td>
</tr>
<tr>
<td>2002</td>
<td>6.7 (45/2.3–17.9)</td>
<td>25.0 (36/13.7–41.1)</td>
</tr>
<tr>
<td>2003</td>
<td>0.0 (2/0.0–65.8)</td>
<td>33.3 (6/9.7–70.0)</td>
</tr>
<tr>
<td>2004</td>
<td>0.0 (1/0.0–79.4)</td>
<td>14.3 (7/2.6–51.3)</td>
</tr>
<tr>
<td>2005</td>
<td>33.3 (3/6.2–79.2)</td>
<td>16.7 (6/3.0–56.4)</td>
</tr>
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A Denotes the naïve population of 1999, the year of WNV introduction, and includes both juvenile and adult crows.

Fig. 1. Estimated naïve American crow (top) and fish crow (bottom) mortality from WNV in New Jersey for the 1999 to 2005 transmission seasons. Naïve populations include both adult and juvenile crows in 1999 (year of WNV introduction) and only juvenile crows for the years 2000–2005. Points and error bars represent the medians and 95% credible intervals.

Fig. 2. Sampling distribution of mortality change (Δm) for each crow species, 1999 to 2005. Each distribution is the standardized frequency distribution of 150,000 Bayesian samples (the area under each curve is equal to 1). For American crow, Δm was negative in 90% of the samples and for fish crow in 60% of the samples.
crow. Thus, American crow should continue to be an efficient surveillance target in areas where populations remain dense and, thus, carcasses are likely to be detected and reported.

REFERENCES


