# Vertical Transmission of Avian Bornavirus in Psittacines

To the Editor: Proventricular dilatation disease (PDD) is a fatal disease in psittacines that jeopardizes critical species conservation projects, such as that involving the Spix's macaw (Cyanopsitta spixii), the world's most endangered bird species (1). The disease is characterized by lymphoplasmacytic infiltrations in the enteric and central nervous systems (2). Consequently, gastrointestinal and neurologic disorders are the major clinical manifestations. Only recently has the cause of the disease been identified by characterization of a newly discovered member of the family Bornaviridae, the avian bornavirus (ABV), which has been detected in affected psittacines (3,4). The relationship of an infection with ABV and the occurrence of PDD has been described in natural cases (5,6) and in experimental trials (7,8). However, birds that are infected with ABV but that are clinically healthy have also been recognized (6). Infected birds can shed viral RNA intermittently (9), and not all infected birds seroconvert (5). For psittacine flock management, control of an ABV infection is critical, e.g., by repeated testing of breeding stock and removal of ABV-positive birds (2,5). However, in breeding projects of rare species, every individual is genetically important and cannot be lost. Therefore, pairing infected, but clinically healthy, birds separately from birds that test negative for the virus might represent an option. For this possibility to be viable, whether vertical transmission of ABV can take place must be further clarified. A study investigating the distribution of ABV in tissues of PDD-positive birds has demonstrated ABV antigen in follicular cells, which may point toward vertical transmission (9).

To investigate vertical transmission of ABV, we examined 30 deadin-shell embryos of various psittacine species that originated from ABVinfected flocks with a history of PDD. First, the eggshell was disinfected and opened at the blunt end by using sterile equipment. The brain and proventriculus of each embryo were analyzed for the presence of ABV RNA by using 2 different real-time reverse transcription PCRs, as described by Honkavuori (4), with the primer pair 1034-1322 and, in case of a negative result, the additional primer set 1367. Sampling, RNA extraction, and PCR were repeated by using the same methods to exclude possible crosscontamination of samples. Afterwards, the complete embryo was placed in 10% buffered formalin, and histopathologic examination and immunohistochemical (IHC) testing were carried out (10) with antibodies directed against the viral phosphoprotein and X protein. If ABV RNA or ABV antigen was demonstrated. crop and cloacal swab specimens and serum of the parents of the positive embryo were immediately taken and used either for ABV RNA detection with the above described PCR or for the detection of specific ABV antibodies by indirect immunofluorescence assay (10). This procedure was chosen because earlier sampling of the parents might have caused breeding interruption, and which eggs of which parents would be available for investigation was not clear

In 2 of the 30 embryos investigated, ABV RNA was detected by using the 1034 PCR. The repeated procedure provided the same results in the same embryos. One embryo was a Major Mitchell cockatoo (Cacatua *leadbeateri*) (cvcle threshold 31.41) and the other a red-crowned Amazon (Amazona *viridigenalis*) (cycle threshold 35.1). None of the investigated embryos demonstrated histopathologic lesions typical of an ABV infection. IHC testing did not show any positive results. However, in the ABV-positive Amazon embryo, an equivocal result was obtained.

The swab specimens of both parents of the Major Mitchell cockatoo tested positive for ABV RNA, but serum did not demonstrate specific ABV antibodies. The crop swab specimen of the female red crowned Amazon was positive for ABV RNA, but serum was negative for ABV antibodies; the male bird tested negative by PCR but demonstrated an ABV-specific antibody titer of 80.

These results highlight the potential risk for vertical transmission of ABV and the conclusion that ABVinfected parents can most likely produce infected offspring. However, test results were positive for only 2 of 30 eggs. Because potentially deadin-shell embryos are usually further incubated by the breeder to ensure embryonic death, the possibility cannot be excluded that ABV RNA was already degraded in some cases. thus causing false-negative results. On the other hand, these eggs might have originated from ABVnegative parents. The quality of the samples might also have caused the questionable IHC results. In PDDaffected, ABV-positive flocks >30% of the birds could be infected (5,6)and the virus is shed intermittently (9). Therefore, pairing ABV-positive birds, incubating their eggs artificially, and raising the chicks separately until they show negative test results, might be an option for breeding projects. However, when vertical transmission occurs (and, if so, its incidence) is unknown

Whether ABV infection of the embryos was the cause of death remains unclear. Even if typical lesions were not detected, the poor quality of the material might have hidden such lesions. However, embryonic infection that does not result in embryonic death is the basic requirement for successful vertical transmission. These preliminary results warrant further studies investigating the possibility of vertical transmission by ABV-infected pairs, especially to minimize the risk for such transmission to endangered species with restricted breeding opportunities.

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### Michael Lierz, Anne Piepenbring, Christiane Herden, Kirstin Oberhäuser, Ursula Heffels-Redmann, and Dirk Enderlein

Author affiliations: Justus Liebig University, Giessen, Germany (M. Lierz, A. Piepenbring, C. Herden, U. Heffels-Redmann, D. Enderlein); and Loro Parque Fundacion, Tenerife, Spain (K. Oberhauser)

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Address for correspondence: Michael Lierz, Clinic for Birds, Reptiles, Amphibians and Fish, Faculty of Veterinary Medicine, Justus Liebig University Giessen, Frankfurter Str 91-93, 35392 Giessen, Germany; email: michael. lierz@vetm0ed.uni-giessen.de



## Novel Astroviruses in Children, Egypt

To the Editor: Human astroviruses (HAstVs) are a common cause of gastroenteritis in children, the elderly, and immunocompromised persons (1). Up to 10% of acute viral gastroenteritis in children and 0.5%-15% of diarrheal outbreaks are attributed to astroviruses (2). Until 2008, eight classical astrovirus serotypes were known to cause human disease; in Egypt, HAstV-1 is the most frequent astrovirus serotype detected (3). Recently, 5 novel astroviruses have been discovered in human fecal samples from patients with diarrhea or acute flaccid paralysis (4-7). Because the prevalence of these virsues in the Middle East is unknown, we screened fecal samples from children with diarrhea residing in Egypt to ascertain the prevalence and diversity of these novel astroviruses.

Fecal samples were collected from a cohort of 364 symptomatic children <5 years of age who had diarrhea and were seeking medical care at Abu Homos Hospital in the Nile Delta of Egypt from September 2006 through September 2007. RNA was extracted from 10% fecal suspensions and reverse transcription PCR for astrovirus was performed as previously described (4-6). Astrovirus consensus primer pair SF0073/SF0076 amplified an  $\approx$ 409-bp region of open reading frame (ORF) 1b, encoding the RNA polymerase gene. PCR-positive samples were then tested by using primer sets Mon269-Mon270 (8) and SF0053-SF0061, amplifying either a 449-bp or a 402-bp product of the ORF2 capsid gene from classical HAstVs (serotypes I-VIII), or astrovirus MLB1, respectively. DNA sequences of PCR products were determined by using Big Dye Terminator Cycle technology (Applied Biosystems, Foster City, CA, USA). Nucleotide sequences were compared with sequences obtained from GenBank. Phylogenies