

Many zoonotic diseases have birds as their natural hosts (Shiina *et al.* 2004; Burt 2005). For example, waterfowl are the natural hosts

require biosafety precautions. Standard sampling and storage during avian influenza surveillance is bound to the availability of nearby deep freezers and transport of samples is subjected to strict regulations. Analysis can only take place in specialised laboratories. These facts make avian influenza research almost impossible if not conducted within the infrastructure of one of the few big collaborative projects. Hence, important contributions from the many smaller ecological projects may be missed (Bin Muzaffar *et al.* 2006; Cromie *et al.* 2006).

Here, a possible solution for this problem is examined: a method to sample, store and analyse potential AIV containing samples. This method does not require immediate deep freezing. The issue of preserving RNA viruses for later analysis (Munster *et al.* 2009) has been addressed several times already in similar fields (Li *et al.* 2004b; Moscoso *et al.* 2005; Ndunguru *et al.* 2005; Perozo *et al.* 2006; Purvis *et al.* 2006; Inoue *et al.* 2007; Nuchprayoon *et al.* 2007; Picard-Meyer *et al.* 2007; Muthukrishnan *et al.* 2008). The so-called FTA cards® (Whatman®) are used to preserve AIV RNA on a dry storage basis. The chemicals in the FTA (Flinders Technology Associates) card render pathogens inactive upon contact (Rogers & Burgoyne 1997) and transport can be arranged safely with only few further biosafety measures to be taken. FTA cards would therefore also be suitable for working with highly pathogenic strains of AIV. Proof of the potential of this principle is given in this short communication. The basis of this method is the isolation of the RNA followed by a one-step RT-PCR. The establishment of these protocols will be

possible in any molecular laboratory, without the need for further biosafety measures. Samples can be mailed by normal postal services. Both sampling and analysis will be available to any molecular ecologist, thereby facilitating further scientific progress. This holds new possibilities for innovative studies in the fields of, for instance, molecular ecology, host-pathogen interactions or ecological immunology.

Methods

Wild Mallard were caught in a duck trap at Ottenby Bird Observatory, Sweden (56°12'N 16°24'E), and cloacal samples were taken for AIV detection. Detailed information about trap d informm24.9(A c4t4.9129 -1.3 TD(

The RNA was eluted into 50 µl elution buffer as provided by the kit. Three 2 mm punches from an untreated FTA card were carried along as negative extraction control; that is, to determine any contamination of the laboratory's tools or devices with AIV material. For RT-PCR detection we used the one-step Access RT-PCR System (Promega) – *i.e.* where reverse transcription into cDNA and PCR amplification is carried out in one tube – following a protocol adjusted from Fouchier *et al.* (2000). Stock solutions of 0.5 µl with 100 mM of the primers M52C and M253R (Fouchier *et al.* 2000) were used in reactions containing 10 µl AMV/Th 5× buffer, 1 µl dNTPs, 7 mM MgSO₄bu xtra3R (Fhier

reported where alternative preservation methods were used (< 200 bp in ethanol; Wang *et al.* 2008).

Since it has recently been shown that RNA fragments of > 700 bp in size can be amplified successfully in other systems (Muthukrishnan *et al.* 2008) we assume that storage of avian influenza samples on FTA cards has the potential to be superior to the ethanol fixation method if primers for larger fragments are used. Some studies tested the sensitivity (*e.g.* RNA quantity) required for detection. They reported the detection of a positive signal even after many-fold dilutions (Perozo *et al.* 2006) or for only 0.1 fg of RNA template (Rogers & Burgoyne 2000), and after storage at ambient temperatures for > 2 weeks. Others claim that RNA on FTA cards is stable even after six months of storage under ambient conditions (Rogers & Burgoyne 2000). Whether these methods are applicable under fieldwork conditions remains to be tested. In natural samples like faeces or oral/cloacal swabs there is also the chance that AIV is present in lower concentrations than tested here. This poses the risk of not detecting an avian influenza infection when there actually is one (*i.e.* a false negative). In particular the effects of storage time and temperature, as well as sensitivity at lower concentrations and contamination through faecal material, would need attention in such a systematic test. Recent studies have however shown that PCR is more sensitive than traditional methods, even when AIVs are only present as unviable particles (Runstadler *et al.* 2007). This also makes detection possible when infection is almost cleared by immune response. To this point,

cloacal samples were not tested directly in the present study but it seems that the use of FTA cards in large scale AIV sampling may be the means by which the field of AIV ecology can be lifted beyond the constraints of difficult sampling, storage and laboratory facilities.

Acknowledgements

Technical assistance was provided by Bert Dibbits and Haisheng Nie, and we thank Jan van der Poel for helpful discussions on preparing for the experiment. Further we would like to thank the Animal Breeding and Genomics Group, Wageningen University, Wageningen, The Netherlands, for hosting us in their laboratory and the Ottenby Bird Observatory, Sweden, for hosting collaborators and providing samples. Financial support was given by the KNJV (Dutch hunters association), the Dutch Ministry of Agriculture, the Faunafonds and the Stichting de Eik Trusts (both in The Netherlands), the Swedish Research Council (grant no. 2007-20774) and the EC-funded Newflubird project. This is contribution No. 230 from the Ottenby Bird Observatory.

References

- Aymard-Henry, M., Coleman, M.T., Dowdle, W.R., Laver, W.G., Schild, G.C. & Webster, R.G. 1973. Influenzavirus neuraminidase and neuraminidase-inhibition test procedures. *Bulletin of the World Health Organization* 48: 199–202.
- Benson, D.A., Karsch-Mizrachi, I., Lipman, D.J., Ostell, J. & Sayers, E.W. 2009. GenBank. *Nucleic Acids Research* 37 (Supplement 1): D26.
- Bin Muzaffar, S., Ydenberg, R.C. & Jones, I.L. 2006. Avian influenza: An ecological and evolutionary perspective for waterbird scientists. *Waterbirds* 29: 243–257.
- Burgoyne, L.A. 1996. Solid medium and method for DNA storage. US Patent No. 5,496,562.

- US Patent and Trademark Office, Washington DC, USA.
- Burt, D.W. 2005. Chicken genome: Current status and future opportunities. *Genome Research* 15: 1692–1698.
- Cavanagh, D. 2005. Coronaviruses in poultry and other birds. *Avian Pathology* 34: 439–448.
- Chen, H., Smith, G.J.D., Li, K.S., Wang, J., Fan, X.H., Rayner, J.M., Vijaykrishna, D., Zhang, J.X., Zhang, L.J., Guo, C.T., Cheung, C.L., Xu, K.M., Duan, L., Huang, K., Qin, K., Leung, Y.H.C., Wu, W.L., Lu, H.R., Chen, Y., Xia, N.S., Naipospos, T.S.P., Yuen, K.Y., Hassan, S.S., Bahri, S., Nguyen, T.D., Webster, R.G., Peiris, J.S.M. & Guan, Y. 2006. Establishment of multiple sublineages of H5N1 influenza virus in Asia: Implications for pandemic control. *Proceedings of the National Academy of Sciences of the United States Of America* 103: 2845–2850.
- Cromie, R.L., Lee, R. & Hughes, B. 2006. Avian influenza: A short review of the disease in wild birds, and of European wild bird surveillance during winter 2005/06. *Wildfowl* 56: 197–202.
- Feare, C.J. & Yasue, M. 2006. Asymptomatic infection with highly pathogenic avian influenza H5N1 in wild birds: how sound is the evidence? *Virology Journal* 3: 96–99.
- Fouchier, R.A.M., Bestebroer, T.M., Herfst, S., Van der Kemp, L., Rimmelzwaan, G.F. & Osterhaus, A. 2000. Detection of influenza A viruses from different species by PCR amplification of conserved sequences in the matrix gene. *Journal of Clinical Microbiology* 38: 4096–4101.
- Fouchier, R.A.M., Munster, V., Wallensten, A., Bestebroer, T.M., Herfst, S., Smith, D., Rimmelzwaan, G.F., Olsen, B. & Osterhaus, A. 2005. Characterization of a novel influenza A virus hemagglutinin subtype (H16) obtained from black-headed gulls. *Journal of Virology* 79: 2814–2822.
- Gilbert, M., Xiao, X.M., Domenech, J., Lubroth, J., Martin, V. & Slingenbergh, J. 2006. Anatidae migration in the western palearctic and spread of highly pathogenic avian influenza H5N1 virus. *Emerging Infectious Diseases* 12: 1650–1656.
- Hoffmann, E., Stech, J., Guan, Y., Webster, R.G. & Perez, D.R. 2001. Universal primer set for the full-length amplification of all influenza A viruses. *Archives of Virology* 146: 2275–2289.
- Inoue, R., Tsukahara, T., Sunaba, C., Itoh, M. & Ushida, K. 2007. Simple and rapid detection of the porcine reproductive and respiratory syndrome virus from pig whole blood using filter paper. *Journal of Virological Methods* 141: 102.
- Kida, H., Yanagawa, R. & Matsuoka, Y. 1980. Duck influenza lacking evidence of disease signs and immune-response. *Infection and Immunity* 30: 547–553.
- Koehler, A.V., Pearce, J.M., Flint, P.L., Franson, J.C. & Ip, H.S. 2008. Genetic evidence of intercontinental movement of avian influenza in a migratory bird: The northern pintail (*Anas acuta*). *Molecular Ecology* 17: 4754–4762.
- Latorre-Margalef, N., Gunnarsson, G., Munster, V.J., Fouchier, R.A.M., Osterhaus, A.D.M.E., Elmberg, J., Olsen, B., Wallensten, A., Haemig, P.D., Fransson, T., Brudin, L. & Waldenström, J. 2009. Effects of influenza A virus infection on migrating mallard ducks. *Proceedings of the Royal Society B: Biological Sciences* 276: 1029.
- Li, K. S., Guan, Y., Wang, J., Smith, G.J.D., Xu, K.M., Duan, L., Rahardjo, A.P., Puthavathana, P., Buranathai, C., Nguyen, T.D., Estoepongastie, A.T.S., Chaisingh, A., Auewarakul, P., Long, H.T., Hanh, N.T.H., Webby, R.J., Poon, L.L.M., Chen, H., Shortridge, K.F., Yuen, K.Y., Webster, R.G. & Peiris, J.S.M. 2004a. Genesis of a highly pathogenic and potentially pandemic H5N1

- influenza virus in eastern Asia. *Nature* 430: 209–213.
- Li, C.C., Beck, I.A., Seidel, K.D. & Frenkel, L.M. 2004b. Persistence of human immunodeficiency virus type 1 subtype B DNA in dried-blood samples on FTA filter paper. *Journal of Clinical Microbiology* 42: 3847.
- MoscOSO, H., Raybon, E.O., Thayer, S.G. & Hofacre, C.L. 2005. Molecular detection and serotyping of infectious bronchitis virus from FTA® filter paper. *Avian Diseases* 49: 24.
- Munster, V.J., Baas, C., Lexmond, P., Waldenström, J., Wallensten, A., Fransson, T., Rimmelzwaan, G.F., Beyer, W.E.P., Schutten, M., Olsen, B., Osterhaus, A. & Fouchier, R.A.M. 2007. Spatial, temporal, and species variation in prevalence of influenza A viruses in wild migratory birds. *PLoS Pathogens* 3: 630–638.
- Munster, V.J., Baas, C., Lexmond, P., Bestebroer, T.M., Guldemeester, J., Beyer, W.E.P., De Wit, E., Schutten, M., Rimmelzwaan, G.F., Osterhaus, A.D.M.E. & Fouchier, R.A.M. 2009. Practical considerations for high-throughput influenza A virus surveillance studies of wild birds by use of molecular diagnostic tests. *Journal of Clinical Microbiology* 47: 666.
- Muthukrishnan, M., Singanallur, N.B., Ralla, K. & Villuppanoor, S.A. 2008. Evaluation of FTA® cards as a laboratory and field sampling device for the detection of foot-and-mouth disease virus and serotyping by RT-PCR and real-time RT-PCR. *Journal of Virological Methods* 151: 311.
- Ndunguru, J., Taylor, N.J., Yadav, J., Aly, H., Legg, J.P., Aveling, T., Thompson, G. & Fauquet, C.M. 2005. Application of FTA technology for sampling, recovery and molecular characterization of viral pathogens and virus-derived transgenes from plant tissues. *Virology Journal* 2: 45.
- Nuchprayoon, S., Saksirisampant, W., Jaijakul, S. & Nuchprayoon, I. 2007. Flinders Technology Associates (FTA) filter paper-based DNA extraction with Polymerase Chain Reaction (PCR) for detection of *Pneumocystis jirovecii* from respiratory specimens of immunocompromised patients. *Journal of Clinical Laboratory Analysis* 21: 382.
- Olsen, B., Munster, V.J., Wallensten, A., Waldenström, J., Osterhaus, A. & Fouchier, R.A.M. 2006. Global patterns of influenza A virus in wild birds. *Science* 312: 384–388.
- Perozo, F., Villegas, P., Estevez, C., Alvarado, I. & Purvis, L.B. 2006. Use of FTA® filter paper for the molecular detection of Newcastle disease virus. *Avian Pathology* 35: 93–98.
- Picard-Meyer, E., Barrat, J. & Cliquet, F. 2007. Use of filter paper (FTA®) technology for sampling, recovery and molecular characterisation of rabies viruses. *Journal of Virological Methods* 140: 174.
- Purvis, L.B., Villegas, P. & Perozo, F. 2006. Evaluation of FTA® paper and phenol for storage, extraction and molecular characterization of infectious bursal disease virus. *Journal of Virological Methods* 138: 66.
- Rimmelzwaan, G.F., Baars, M., Claas, E.C.J. & Osterhaus, A. 1998. Comparison of RNA hybridization, hemagglutination assay, titration of infectious virus and immunofluorescence as methods for monitoring influenza virus replication in vitro. *Journal of Virological Methods* 74: 57–66.
- Rogers, C.D.G. & Burgoyne, L. 1997. Bacterial typing: Storing and processing of stabilized reference bacteria for polymerase chain reaction without preparing DNA – An example of an automatable procedure. *Analytical Biochemistry* 247: 223–227.
- Rogers, C.D.G. & Burgoyne, L.A. 2000. Reverse transcription of an RNA genome from databasing paper (FTA®). *Biotechnology and Applied Biochemistry* 31: 219–224.
- Runstadler, J.A., Happ, G.M., Slemmons, R.D., Sheng, Z.-M., Gundlach, N., Petrus, M.,

- Senne, D., Nolting, J., Evers, D.L., Modrell, A., Huson, H., Hills, S., Rothe, T., Marr, T., Taubenberger, J.K.. 2007. Using RRT-PCR analysis and virus isolation to determine the prevalence of avian influenza virus infections in ducks at Minto Flats State Game Refuge, Alaska, during August 2005. *Archives of Virology* 152: 1901–1910.
- Salk, J.E. 1944. Simplified procedure for titrating hemagglutinating capacity of influenza virus and the corresponding antibody. *Journal of Immunology* 49: 87–98.
- Serratos, J., Ribo, O., Correia, S. & Pittman, M. 2007. EFSA scientific risk assessment on animal health and welfare aspects of avian influenza (EFSA-Q-2004-075). *Avian Diseases* 51: 501–503.
- Shiina, T., Shimizu, S., Hosomichi, K., Kohara, S., Watanabe, S., Hanzawa, K., Beck, S., Kulski, J.K. & Inoko, H. 2004. Comparative genomic analysis of two avian (quail and chicken) MHC regions. *Journal of Immunology* 172: 6751–6763.
- Spackman, E., Senne, D.A., Myers, T.J., Bulaga, L.L., Garber, L.P., Perdue, M.L., Lohman, K., Daum, L.T. & Suarez, D.L. 2002. Development of a real-time reverse transcriptase PCR assay for type A influenza virus and the avian H5 and H7 hemagglutinin subtypes. *Journal of Clinical Microbiology* 40: 3256.
- van Gils, J.A., Munster, V.J., Radersma, R., Liefhebber, D., Fouchier, R.A.M. & Klaassen, M. 2007. Hampered foraging and migratory performance in swans infected with low-pathogenic Avian Influenza A Virus. *PLoS ONE* 2: e184.
- Wallensten, A., Munster, V.J., Karlsson, M., Lundkvist, A., Brytting, M., Stervander, M., Osterhaus, A., Fouchier, R.A.M. & Olsen, B. 2006. High prevalence of influenza A virus in ducks caught during spring migration through Sweden. *Vaccine* 24: 6734–6735.
- Wallensten, A., Munster, V.J., Latorre-Margalef, N., Brytting, M., Elmberg, J., Fouchier, R.A.M., Fransson, T., Haemig, P.D., Karlsson, M., Lundkvist, A., Osterhaus, A., Stervander, M., Waldenström, J. & Olsen, B. 2007. Surveillance of influenza A virus in migratory waterfowl in northern Europe. *Emerging Infectious Diseases* 13: 404–411.
- Wang, R., Soll, L., Dugan, V., Runstadler, J.A., Happ, G., Slemons, R.D. & Taubenberger, J.K. 2008. Examining the hemagglutinin subtype diversity among wild duck-origin influenza A viruses using ethanol-fixed cloacal swabs and a novel RT-PCR method. *Virology* 375: 182.
- Weber, T.P. & Stilianakis, N.I. 2007. Ecologic immunology of avian influenza (H5N1) in migratory birds. *Emerging Infectious Diseases* 13: 1139–1143.
- Webster, R.G., Bean, W.J., Gorman, O.T., ChamberTJ-10.2275 -Ko.a0AlasY-36.9(y)TJean,3g-