

Identification of FPV by PCR amplification of p4b gene in infected cell culture and Chorioallantoic membrane

R.K. Verma¹, R.K. Joshi² and Shubhra Shukla³

¹Department of Veterinary Teaching Veterinary Clinical Complex (T.V.C.C), College of Veterinary Science and Animal Husbandry N.D. University of agriculture and Technology, Kumarganj- 224229, Faizabad (U.P.), India.

²Department of Veterinary Microbiology, College of Veterinary Science and Animal Husbandry N.D. University of agriculture and Technology Kumarganj- 224229, Faizabad (U.P.), India.

³Department of Biosciences, Integral University, Lucknow (U.P.) 226026, India

(Received: 20 May 2015; accepted: 11 August 2015)

Five local fowl were brought for postmortem examination with a history of sudden death in the area of Kumarganj, Faizabad (U.P.). The gross examination of birds revealed multiple light whitish nodules around the eye, on the skin at the level of hock joint, on the anterior part of tracheal mucosa, congested lung and pallor liver. Impression smears from nodules revealed numerous heterophils, red blood cells, necrotic epithelial cells and bacterial colonies. Histopathological examination of nodules revealed eosinophilic intracytoplasmic inclusions (Bollinger bodies) in keratinocytes, epidermal hyperplasia and necrosis with ballooning degeneration, and bacterial colonies. The virus was isolated and infection was produced on both chorioallantoic membrane and BGM-70. Polymerase chain reaction was carried out and primer set designed from the 4b core protein gene of fowl poxvirus revealed amplification at 576bp.

Key words: Avianpox virus, Polymerase chain reaction (PCR), BGM-70, Chorioallantoic membrane.

The largest known virus of terrestrial and marine mammals belongs to the *Chordopoxvirinae* subfamily of the *Poxviridae* family¹. The Poxviridae is divided into two subfamilies: Entomopoxvirinae, that comprises insect poxviruses, and Chordopoxvirinae, that includes all poxviruses of vertebrates². The family Poxviridae possess non-infectious, double stranded DNA genomes that range in size from 130-380 kbp and replicate almost exclusively in the cell cytoplasm². Poxviruses are highly adapted viruses infecting a large number of hosts, including insects, reptiles, birds, over 30 mammalian species and in a number of endangered species or species in captive breeding recovery programs^{3,4}. Latest case report of Fowlpox virus

infection is found in Hungarian partridges (*Perdix perdix*)⁵, white-tailed sea eagle (*Haliaeetus albicilla*)⁶, and Hungarian great tits (*Parus major*)⁷. APVs are transmitted via biting insects and aerosols and are usually named virus was first isolated and characterized⁸. Interspecies transmission also occurs of fowlpox virus⁹. The avipox virus appears incapable of causing disease in mammals; therefore there appears to be no zoonotic potential¹⁰. The disease characterized by proliferative lesions of the skin and diphtheric membranes of the respiratory tract, mouth and esophagus, has been described in avian species⁸. Large intracytoplasmic inclusion bodies known as Bollinger bodies that contain smaller elementary bodies (Borrel bodies) is formed when FPV multiplies in cytoplasm of epithelium is characteristic of the disease and can be seen in histopathological examination¹¹.

* To whom all correspondence should be addressed.
Tel.: +919369382948;
E-mail: drrajesh16@yahoo.com

APV are increasingly being detected and characterized by PCR, Restriction fragment length polymorphism (RFLP), Southern blot hybridization, and cycle sequencing, directed at specific genes such as the 4b core protein gene^{12,13}. Prior to PCR technology, viruses were isolated in cell culture before any further analyses could occur¹⁴. Identification and differentiation of viruses before the implementation of PCR relied on less specific serologic assays such as, virus neutralization, hemagglutination and immuno fluorescence assays, later complemented by restriction endonuclease profiles of viral DNA resolved in agarose or poly acrylamide gels^{15,16,17,18,2}. A recombinant fowlpox virus vaccine expressing key protective *Mycoplasma gallisepticum* antigens could facilitate in the prevention both of fowlpox virus and *M. gallisepticum* infections¹⁹. Pigeon pox virus may be used as first dose in chickens against fowl pox. This is preferred especially for layers to avoid any reaction *Ganguli*²⁰.

Considering poxviruses in particular, PCR and genome sequencing has meant the evolution from sometimes vague histopathologic and electron microscopic (EM) diagnoses to much more definitive genetic assays for poxvirus infection^{21,22,23}. Naturally occurring dual infection of layer chickens with fowlpox virus and gallid herpesvirus 1 (infectious laryngotracheitis virus) can be diagnosed with the help of electron microscopy, PCR and histology²⁴. Because of this common poxvirus morphology, it is difficult to discern between genera of poxviruses when using techniques like histopathology and EM. However, PCR and sequencing methods reveal not only the genus, but in most cases, species of the virus being examined^{21,25,23}. The advent of PCR allowed for the direct amplification of viral DNA and rapid genome sequencing². These sequencing advances have allowed for a better ability to define and understand the evolutionary relationships between the different poxvirus genera. Comparing genes that have been identified as highly conserved can aid in new virus characterization and comparison. Previous studies have elected phylogenetic analysis to be the best tool available for characterizing poxviruses known to date^{26,25,27}. PCR allows for sensitive and specific detection of viral nucleic acids and has been shown to increase the diagnostic sensitivity for many viral pathogens when compared to culture. Detection

by realtime PCR has been used to identify recombinant APV from individual plaques²⁸.

MATERIALS AND METHOD

Detection of FPV using polymerase chain reaction

The FPV specific viral DNA was amplified and detected in FPV infected CAM as well as BGM-70 cell solution samples.

Extraction of viral DNA

The method described by Eo.²⁹ was used for extraction of viral DNA from FPV infected CAM suspensions.

The FPV infected CAM was triturated in TE buffer (10 mM Tris and 1 mM EDTA, pH 8.0) and centrifuged at 150xg at 4°C for 10 min. In 2 ml appendroff tube, to 400µl suspension of each sample, equal volume of extraction buffer was added and tubes were incubated at 55°C in water Bath for 2 hr. Equal volume of Ph:Cl (48:2) solution was then added to the mixture and tubes were vortexed followed by centrifugation at 13000 rpm for 10 min. The aqueous phase was collected in another appendroff tube and the step was repeated once. Following centrifugation at 13000 rpm for 10 min, the aqueous phase was collected and to this, equal volume of chloroform was added and tubes were vortexed. The suspension was then centrifuged at 13000 rpm for 10 min and aqueous phase was collected. The step was repeated once with chloroform and to the aqueous phase, 1/10 volume of 3M Sodium Acetate and 2 volume of absolute ethanol was added. The mixture was kept overnight at -70°C. The mixture was then centrifuged at 13000 rpm for 30 min at 4°C, supernatant was removed and again 700µl of 70% ethanol was added, mixed properly and centrifuged at 13000 rpm for 10 min. After removing the supernatant, tubes were allowed to dry. To the product, 50µl of TE buffer was added after drying the tube and it was stored at -20°C till further use.

Primers

PCR was performed to amplify the p4b core protein gene of avian pox virus. The published primers, forward (CP1, 5'-CAGCAGGTGCTAA CAACAA-3') and reverse (CP2, 5'-CGGTAGCTTAACGCCGAATA-3') specific to the p4b core protein gene (Weil¹³ et al., 2004) of avian pox virus were used

Composition of Reaction mixture for PCR

Reaction mixture was prepared in 50 μ l volume for one PCR reaction as given below:

Table 1.

Master Mix (10mM)	25.0 μ l
Forward Primer (50.0 picomol)	5.0 μ l
Reverse Primer (50.0 picomol)	5.0 μ l
Nuclease free water	10 μ l
DNA Template	5.0 μ l
Total	50.0 μ l

Steps and conditions of thermal cycling

Following cycling conditions were used as described by Eo²⁹

Table 2.

Parameters	Temperature	Time
Heat lid	105°C	
Initial denaturation	94°C	5 minutes
No. of cycle	35	
Denaturation	94°C	20 second
Annealing	53°C	30 second
Extension	72°C	40 second
Final Extension	72°C	5 minutes
Final hold	04°C	

Confirmation of PCR Products

The amplified PCR products were confirmed for their expected size in 1.5% agarose gel in 0.5X TBE buffer using horizontal submarine electrophoresis apparatus. A 1.5 percent agarose gel prepared in 0.5X TBE buffer was boiled for 2 min and allowed to cool down to 50°C, Ethidium bromide was added @ 2 μ l/50 ml and mix thoroughly. The gel casting platform was placed on a leveled surface and the open slides were sealed with adhesive tape. The gel comb was then placed across the gel casting platform, so that the teeth of comb remained 1 mm above the base of platform. The molten agarose was then poured onto the gel casting platform and it was kept undisturbed until the gel was properly solidified. The comb was taken out and adhesive tape was removed. The set gel with gel casting platform was then submerged in the sufficient quantity (about 1 mm above the gel level) of electrophoresis buffer keeping the wells towards cathode end.

Eight μ l of PCR products and molecular weight markers (100bp DNA ladder) were mixed with 2 μ l of Bromophenol blue dye (6X) and loaded into

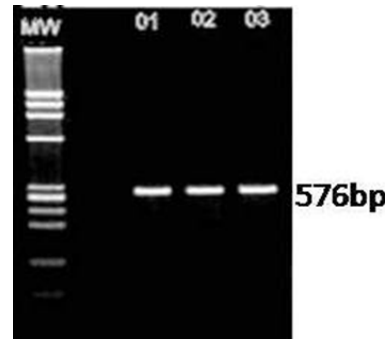
the respective wells. Electrophoresis was performed at 5 volt/cm current and the progress of motility was monitored by the migration of dye. At the end of electrophoresis, the gel was visualized under Gel documentation system (UV Tech.) for the amplified product of desired length.

RESULTS

On PCR analysis using P4b gene specific primer set, the P4b gene could be amplified in both CAM as well as BGM-70 cell infected with the local isolate of the FPV with an amplification size of 576 bps.

DISCUSSION

In present study P4b gene was successfully amplified in the CAM as well as BGM-70 cells infected with the field isolate of FPV with an amplicon size of 576 bps. PCR and sequencing



Lane-MW: 100 bp DNA ladder,
Lane 01: FPV vaccine virus
Lane 02: CAM grown FPV
Lane 3: Cell culture grown virus.
PCR amplification of P4b core protein gene



Fig. 1. Fowl Pox Virus in Avian.

are reported to be the methods of choice for rapid identification and differentiation of poxviruses up to the species level^{30,31,32}. The P4b gene was targeted by many workers for identification of Avipox viruses in infected tissue because P4b is a conserved 752 kDa virion core protein found in all poxviruses^{33,34}. Weil and Eo^{13,29} also used PCR to amplify the P4b core protein gene to identify avian pox virus and could get a 576 bp product. Williams³⁵ also used PCR on nucleic acid extracted from Vectormune FP-LT (fowl pox virus-laryngotracheitis) vaccine strain and a live attenuated fowl pox vaccine and from paraffin embedded lung tissues from FPV-LT vaccinated birds. PCR-based diagnosis is considered superior to other techniques in terms of sensitivity and speed³⁰. PCR amplification of the 4b core poxvirus gene sequence was successful in four of five individuals resulting in products that averaged 575 bp³⁶. Polymerase chain reaction was carried out and primer set designed from the 4b core protein gene of fowl pox virus revealed amplification at 578 bp³⁷.

Summary

On PCR analysis using P4b gene specific set, the P4b gene could be amplified in both CAM as well as BGM-70 infected with the local isolate of FPV with an amplification size of 576bps.

ACKNOWLEDGMENT

The authors are thankful to Dean, College of Veterinary Science and Animal Husbandry, NDUAT, Kumarganj, Faizabad for providing the funding and facilities to carry out the work.

REFERENCES

1. Fauquet, C. M., Mayo, M. A., Maniloff, J., Desselberger, U. and Ball, L.A. Virus Taxonomy: VIIIth Report of the International Committee on Taxonomy of Viruses". Elsevier Academic Press. 2005.
2. Moss, B. Poxviruses. In Fields Virology, 4th ed., Vol.2, D. M. Knipe, P. M. Howley, D. E. Griffin, R. A. Lamb, M. A. Martin, B. Roizman and S. E. Straus (eds.) Lippincott Williams & Wilkins publishers, Philadelphia, Pennsylvania. 2001; 2849-2884.
3. Bohls, R. L., Linares, J. A., Gross, S. L., Ferro, P. J., Silvy, N. J. and Collisson, E. W. Phylogenetic analyses indicate little variation among reticuloendotheliosis viruses infecting avian species, including the endangered Attwater's prairie chicken. *Virus Res.*, 2006; **119**:187-94.
4. Upton, C., Slack, S., Hunter, A. L., Ehlers, A., and Roper, R. L. 2003. Poxvirus orthologous clusters: toward defining the minimum essential poxvirus genome. *Journal of Virology.*, 2003; **77**(13): 7590-7600.
5. Brower, A.I., Cigel, F., Radi, C., Toohey-Kurth, K. Beak necrosis in Hungarian partridges (*Perdix perdix*) associated with beak-bits and avian poxvirus infection. *Avian Pathol.*, 2010; **39**: 223-5.
6. Saito K, Kodama A, Yamaguchi T, Gotoh Y, Sakai H, Fukushi H, Masegi T, Yanai T. Avian poxvirus infection in a White-tailed Sea Eagle (*Haliaeetus albicilla*) in Japan. *Avian Pathol.*, 2009; **38**: 485-489
7. Palade E.A, Biro N, Dobos-Kovács M, Demeter Z, Mándoki M, Rusvai M: Poxvirus infection in Hungarian great tits (*Parus major*): case report. *Acta. Vet. Hung.*, 2008, **56**: 539-46.
8. Tripathy, D. N., Schnitzlein W. M., Morris, P. J., Janssen D. L., Zuba, J. K., Massey, G. and Atkinson, T. (2000). Characterization of poxviruses from forest birds in Hawaii. *J. Wild L. Dis.*, 2000; **36**: 225-30.
9. Terasaki, T., Kaneko, M., Mase, M. Avian poxvirus infection in flamingos (*Phoenicopterus roseus*) in a zoo in Japan. *Avian Dis.*, 2010; **54**: 955-7.
10. Jarmin, S., Ruth, M., Richard, E.G., Stephen, M.L., Michael, A.S. Avian poxvirus Phylogenetics: Identification of a PCR length polymorphism that discriminates between the two major clades. *Journal of General Virology, Central Veterinary Laboratories: Weybridge, UK.* 2006; **87**: 2191-2201.
11. Alehegn, E., Chanie, M., Desalegne, Mengesha, D. A Systematic Review of Serological and Clinicopathological Features and Associated Risk Factors of Avian Pox. *Brit. J. Poult Sci.*, 2014; **3**(3): 78-87.
12. Luschow, D., Hoffmann, T. and Hafez, H. M. Differentiation of avian poxvirus strains on the basis of nucleotide sequences of 4b gene fragment. *Avian Dis.*, 2004; **48**: 453-462.
13. Weli, S. C., Traavik, T., Tryland, M., Coucheron, D. H. and Nilssen Ø. Analysis and comparison of the 4b core protein gene of avipoxviruses from wild birds: Evidence for interspecies spatial phylogenetic variation. *Arch Virol.*, 2004; **149**: 2035-2046.
14. Levine, A. J. The origins of Virology. In *Fundamental Virology*, 4th ed., D. M. Knipe, P.

- M. Howley, D. E. Griffin, R. A. Lamb, M. A. Martin, B. Roizman and S. E. Straus (eds.) Lippincott, Williams & Wilkins Publishers, Philadelphia, Pennsylvania. 2001; pp.3 -19.
15. Fleming, S. B., Blok, J., Fraser, K. M., Mercer, A. A. & Robinson, A. J. Conservation of gene structure and arrangement between vaccinia virus and orf virus. *Virology.*, 1993; **195**: 175-184.
 16. Robinson, A. J., and Mercer, A., A. Parapox of Red Deer: Evidence for its inclusion as a new member in the genus parapoxvirus. *Virology.*, 1995; **208**: 812-815.
 17. Ropp, S. L., Jin, Q., Knight J. C., Massung, R. F., and Esposito, J. J. PCR Strategy for identification and differentiation of small pox and other orthopoxviruses. *Journal of Clinical Microbiology.*, 1995; **33**(8): 2069-2076.
 18. Mangana-Vougiouka, Odour., Markoulatos, P., Koptopoulos, G., Nomikou, K., Bakandritsos, N., and Papadopoulos, O. Sheep poxvirus identification by PCR in cell cultures. *Journal of Virological Methods.*, 1999; **77**:75-79.
 19. Zhang, G.Z., Zhang, R., Zhao, H.L., Wang, X.T., Zhang, S.P., Li, X.J., Qin, C.Z., Lv, C.M., Zhao, J.X., Zhou, J.F. A safety assessment of a fowlpox vectored *Mycoplasma gallisepticum* vaccine in chickens. *Poult Sci.*, 2010; **89**: 1301-6.
 20. Ganguli, S. Developing Trends in Livestock and Poultry Vaccines: *A Rev Res J. Ani. Vet. Fish. Sci.*, 2013; **1**(3), 25-27.
 21. Damaso, C. R., Esposito, J. J., Condit, R. C., Moussatche, N. (2000). An emergent Poxvirus from humans and cattle in Rio de Jenerio State: Contagalovirus may device from Brazilian small pox vaccine. *Virology.*, 2000; **227**: 439-449.
 22. Gubser, C. and Smith, G. L. The sequence of camelpox virus shows it is most closely related to variola virus, the cause of smallpox. *J Gen Virol.*, 2002; **83**: 855-872.
 23. Hosamani, M., Mondal, B., Tembhurna, P. A., Bandyopadhyay, S. K., Singh, R. K., and Rasool, T. J. Differentiation of sheep pox and goat poxviruses by sequence analysis and PCR-RFLP of P32 gene. *Virus Genes.*, 2004; **29**(1): 73-80
 24. Diallo, I. S., Taylor, J., Gibson, J., Hoad, J., De Jong, A., Hewitson, G., Corney, B.G. Rodwell, B.J. Diagnosis of a naturally occurring dual infection of layer chickens with fowlpox virus and gallidherpesvirus 1 (infectious laryngotracheitis virus). *Avian Pathol.*, 2010; **39**: 25-30.
 25. Becher, P., Konig, M., Muller, G., Siebert, U. & Thiel, H. J. Characterization of sealpox virus, a separate member of the parapoxviruses. *Arch Virol.*, 2002; **147**: 1133-1140.
 26. Afonso, C. L., Tulman, E. R., Lu, Z., Zsak, L., Kutish, G. F. and Rock, D. L. The genome of fowlpox virus. *J Virol.*, 2000; **74**: 3815-3831.
 27. Gubser C, Hue S, Kellam P, and Smith GL: Poxvirus genomes: a phylogenetic analysis. *J Gen Virol.*, 2004; **85**: 105-117.
 28. Boyle D. B., Anderson M. A., Amos R., Voysey R., Coupar B. E. Construction of recombinant fowlpox viruses carrying multiple vaccine antigens and immunomodulatory molecules. *Biotechniques.*, 2004; **37**:104-6, 108-11.
 29. Eo, K. Y., Kim, Y. H., Cho, K. H., Jang, J. S., Kim, T. H., Kwak, D. and Kwon, O. D. Infection of avian pox virus in oriental turtle-doves. *Pakistan Vet. J.*, 2011; **31**(4): 354-356.
 30. Mercer, A.A., Schmidt, A. and Weber, O. Poxviruses (Birkhäuser Advances in Infectious Diseases). Birkhauser, Verlag, Basel. 2007: 441.
 31. Balamurugan, V., Bhanuprakash, V., Hosamani, M., Jayappa, K. D., Venkatesan, G., Chauhan, B. and Singh, R.K. A polymerase chain reaction strategy for the diagnosis of camelpox. *J. Vet. Diagn. Invest.*, 2009; **21**: 231-237.
 32. Yousif, A. A., Al-Naeem, A. A. and Al-Ali, M. A. Rapid non-enzymatic extraction method for isolating PCR-quality camelpox virus DNA from skin. *J. of Virol. Methods.*, 2010; **169**: 138-142.
 33. Binns, M. M., Bournsnel, M. E. G., Tomley, F. M. and Compbell, J. (1989). Analysis of the fowl pox virus gene encoding the 4b core polypeptide and demonstration that it possesses efficient promoter sequence. *J. Virol.*, 1989; **170**: 288-291.
 34. Huw Lee L. and Hwa Lee K. Application of the polymerase chain reaction for the diagnosis of fowl poxvirus infection. *J Virol Methods.*, 1997; **63**:113-9.
 35. Williams, S. M., Smith, J. A., Garcia, M., Brinson, D., Kiupel, M. and Hofacre, C. Severe histiolympocytic and heterophilic bronchopneumonia as a vaccination in broiler chicks reaction to in ovofowlpox. *Vet. Pathol.*, 2010; **47**: 177.
 36. Loreto A. Godoy, Lisa S. Dalbeck, Lisa A. Tell, Leslie W. Woods, Rita R. Colwell, Barbara Robinson, Susan M. Wethington, Anneke Moresco, Peter R. Woolcock, and Holly B. Ernest Characterization Of Avian Poxvirus In Anna's summingbird (*Calypteanna*) In California, USA. *Journal of Wildlife Diseases.*, 2013; **49**(4): 978-985.
 37. Balachandran, C. Pazhanivel, N. Prabhakar, T.G. Murugadas, V. Prabakar P. Avipox Virus Infection in *Rosella parakeet* (*Platycercus* sp.) *Journal of Advanced Veterinary Research.*, 2012; **2**: 184-187.