Diagnosis and histopathological study of avian influenza virus-H5 (AIV-H5) in broiler farms

F.A. Isihak¹, H.Kh. Ismail² and A.A. Wahid³

¹ Department of Microbiology, ² Department of Pathology and Poultry Diseases, College of Veterinary Medicine, University of Mosul, ³General Manager, Yarabarz Company for Poultry Feeding and Veterinary Services, Erbil, Iraq Email: ¹fanar1976@yahoo.com, ²hana_khismail@yahoo.com, ³abdulwahid.abdulhakeem@gmail.com

(Received April 29, 2019; Accepted June 24, 2019)

Abstract

This study was conducted for diagnosis and description of the pathological changes of AIV-H5 as the causative pathogen in Iraqi broiler farms. The current study was carried out on 84 broiler farms. Infected birds were tested for detection of the AIV infection from the tracheal swabs by rapid chromatographic AIV type A and H5 test kits. In RRT-PCR 8 samples (8 farms) of Trachea were selected to be tested by this assay. Samples of trachea, lung, and spleen from the dead birds with natural AIV-H5 infection were submitted for histopathological examination. seventy-two out of 84 farms tested for AIV-Type A gave positive results, and 58 out of 72 positives for type A-AIV gave a positive result for H5 antigen in a rapid chromatographic strip. The main gross lesions in the trachea of infected birds were severe congestion and hemorrhage. In the RRT-PCR assay, 8 out of 8 samples gave a distinct positive result for this test. The microscopic histopathological examination of infected tracheas showed obvious desquamation of lining epithelium with complete loss of cilia associated with congestion. There was deposition of fibrinous material in the splenic tissue associated with the disappearance of the germinal centers. Thus, we concluded that AIV-H5 infection causes severe pathological and histopathological changes as a result of systemic infection. The RRT-PCR assay was highly sensitive and specific for the detection of highly pathogenic avian influenza virus subtypes.

Keywords: Avian influenza, RRT-PCR, Histopathology, Rapid test, Broilers Available online at <u>http://www.vetmedmosul.com</u>, © 2020, College of Veterinary Medicine, University of Mosul. This is an open access article under the CC BY 4.0 license (<u>http://creativecommons.org/licenses/by/4.0/</u>).

دراسة تشخيصية ونسجية مرضية لفايروس إنفلونزا الطيور النمط - ٥ في حقول فروج اللحم

فنار ابلحد اسحق'، هناء خليل اسماعيل' و عبد الواحد عبد الحكيم واحد"

' فرع الأحياء المجهرية، ` فرع الأمراض وأمراض الدواجن، كلية الطب البيطري، جامعة الموصل، " المدير العام، شركة يارا برز للدواجن والخدمات البيطرية، أربيل، العراق

الخلاصة

لقد أجريت هذه الدراسة لتشخيص ووصف التغيرات المرضية المصاحبة للإصابة بفيروس إنفلونزا الطيور - النمط H5 كمسبب مرضي لحقول الفروج اللاحم في العراق. شملت الدراسة ٨٤ حقل للفروج. تم اختبار الطيور المصابة للكشف عن الخمج بفيروس إنفلونزا الطيور في مسحات القصبة الهوائية عن طريق اختبار الاستشراب المناعي السريع (للنمط A و H5). تم أجراء اختبار تفاعل البلمرة المتسلسل ذي الوقت الحقيقي على ٨ عينات من القصبة الهوائية لثماني من هذه الحقول، كما تم إجراء الفحص النسجي المرضي لعينات القصبة الهوائية والرئة والطحال من الطيور النافقة والمصابة طبيعيا بالنمط H5. أظهرت النتائج أن ٢٧ من ٨٤ حقل تم اختبار ها لمحص الاستشراب المناعي السريع نوع A أعطت نتائج إيجابية و ٥٩ من ٢٢ حقل كان موجباً للنوع H5 وتميزت الأفات الرئيسية في القصبة الهوائية المصابة بالاحتقان الشديد والنزف. وفي اختبار تفاعل البلمرة المتسلسل ذي الوقت الحقيقي للكشف عن عنه عنه م والعوائية المصابة بالاحتقان الشديد والنزف. وفي اختبار تفاعل البلمرة المتسلسل ذي الوقت الحقيقي للكشف عن الرئيسية في القصبة الهوائية المصابة بالاحتقان الشديد والنزف. وفي اختبار تفاعل البلمرة المتسلسل ذي الوقت الحقيقي الرئيسية في العرب والعرب المناعي المربع نوع A أعطت نتائج إيجابية و ٢٨ من ٢٢ حقل كان موجباً للنوع H5. وتميزت الأفات الرئيسية في القصبة الهوائية المصابة بالاحتقان الشديد والنزف. وفي اختبار تفاعل البلمرة المتسلسل ذي الوقت الحقيقي للكشف عن فيروس أنفلونزا الطيور نوع H5 - AIV تبين أن ثمانية من مجموع ثماني عينات أظهرت نتائج إيجابية لهذا الاختبار. أظهر الفحص النسجي المجهري لعينات القصبة المصابة وجود انسلاخ لظهارة القصبة الهوائية مع فقدان كامل للأهداب واحتقان الأوعية الدموية. وفي الرئة المصابة لوحظ تحطم الأسناخ واحتقان متعدد البؤر حول الأوعية الدموية. كما كان هناك ترسب من المواد الفبرينية في أنسجة الطحال مع اختفاء المراكز الانتاشية. وبهذا تلخصت نتائج هذه الدراسة إلى أن الإصابة بفيروس إنفلونزا الطيور النمط H5 سبب تغيرات مرضية ونسجية نتيجة للخمج الجهازي. وان الحساسية العالية والسرعة والنوعية التي يتميز بها اختبار تفاعل البلمرة المتسلس ذي الوقت الموت

Introduction

The highly pathogenic avian influenza virus (AIV) subtype H5N1 is considered one of the important pathogens that lead to severe respiratory disease with high morbidity and mortality in poultry industry. The H5N1 virus has firstly recorded in republic of China in 1996 and currently causes infection in worldwide including Asian, African and European countries (1). Since 1996, H5N1 subtypes undergone significant antigenic drift phenomena and these viruses have been classified into different clades according to phylogenetic features depended on hemagglutinin (HA) gene (2). The rapid diagnostic kits for detection of H5 subtypes are necessary for the control of infection with influenza viruses and these kits are affordable in the market for detection of specific nucleoprotein (NP) of this virus. The colloidal test strip technique give a quick results and does not need sophisticated requirements or special expertness, This technique has many benefits such as simplicity, rapidity, low cost, direct results and no expensive materials and equipment's. Pathogenicity of AIVs varies considerably depending upon host species, infective doses and routes of infection (3). Virus cultivation on embryonated eggs is currently performed for tracheal and cloacal samples from avian spp. are considered as standard diagnostic procedure for isolation, detection and subtyping of influenza A viruses (4). Precise and quick diagnostic procedure for the detection of infection is very important, not only for controlling of disease, but also to facilitate early antiviral medications. Molecular diagnosis of AIV-H5 by RRT-PCR is a rapid choice. The results of subtyping of the causative virus may be available within 24 hours. A pair of primers has also recommended by World Health Organization (WHO) for detection of H5 subtype as a laboratory assay (5). Different of RNA viruses has been detected successfully by RRT-PCR assay with hydrolysis probes (6,7). RRT-PCR technique provides advantages of rapidity, viral loading analysis, sensitivity and specificity in comparison with standard protocol of RT-PCR.

This study aimed for identification of AIV-H5 as a causative agent for outbreaks of severe respiratory illness in Iraqi broiler farms during 2018 with description of main pathological changes during the course of the infection with this virus.

Martials and methods

Broiler farms

The study was carried out on 84 broiler farms. All birds in these farms were clinically suffered from respiratory signs, severe depression, nasal discharge, excessive secretions within the lumen of the trachea associated with severe hemorrhages and high mortality rates. A group of infected birds from each farm were tested for detection the AIV infection from the tracheal swabs by rapid chromatographic AIV test kits (type A) (Bionote, Korea) according to the manufacturers' instructions depend on a monoclonal antibody against the viral nucleoprotein (NP). Tested birds which gave Positive results with rapid chromatographic strip type A undergo for detection the AIV-H5 infection with specific rapid chromatographic test kits for H5 antigen (Bionote, Korea).

Tracheal gross Lesions

Tracheas of infected and dead birds suffered clinically from respiratory signs were investigated for the gross lesions as they are the target organs for AIV-H5 virus.

RNA isolation

The isolation of viral RNA from virus containing tissue (trachea) was carried out by using an SV Total RNA Isolation System according to the manufacturer instructions (Promega Corporation, Madison, Wis., catalog # Z3100).

Primers

Two different sets of sequence specific primers were used targeting the hemagglutinin (HA) genes of the H5 subtype which include (8):

Forward primer (5'-ACAAAGCTCTATCAAAAACCCAAC- 3') Reverse primer (5'-TACCCATACCAACCATCTACCAT- 3')

Real-time Reverse Transcriptase/Polymerase Chain Reaction RRT-PCR

Eight tracheal positive samples (for rapid chromatographic strip-H5) were randomly selected from infected birds to be used in this test. briefly the reactions were performed with the GoTaq[®]1- Step RT-qPCR Kit (Promega, USA, catalog # 6021) by using an applied biosystem step one Real-Time PCR system (48 wells, ABI,

USA). The temperature profile consist of an initial step \geq 37 °C for 15 min for reverse transcription process, RT-PCR inactivation/hot-start activation step of 95 °C for 10 min followed by 40 cycles for 10 s at 95 °C for denaturation, 30s at 60 °C for annealing, 30s at 72 °C for elongation and a final extension step at 60-95 °C. The qPCR assay used applied Biosystems SYBR[®] Green universal PCR with master mix (25µl), 5 µl of Forward primer, 5 µl of reverse primer, 10 µl of cDNA produced from reverse transcribed clinical sample, 5 µl MgCl₂ and 5 µl of CXR reference dye.

Histopathology

Samples from the dead birds with natural AIV-H5 infection that detected by rapid chromatographic test were submitted for postmortem and histopathological examination. Tissue section of trachea, lung and spleen were removed and fixed in 10% of neutral buffered formalin. All tissue samples were embedded in paraffin wax, sectioned at 5 μ m, were stained on clean glass slides with Hematoxylin and Eosin (H&E) and Masson trichrome stains for histopathological examination by light microscope (9,10).

Results

Rapid chromatographic strip

Seventy two out of 84 farms tested for AIV were positive (Table 1, Figure 1a) and 58 out of 72 positives for type A-AIV gave positive results for H5 antigen in rapid chromatographic strip (Figure 1b).

Table 1: Percentage and number of tested and infected farms with rapid chromatographic strip

Total number of farms tested for type A	84
Number of Positive AIV-A farms	72
Percentage of Positive AIV-A farms	85.72%
Number of negative AIV-A farms	12
Percentage of negative AIV-A farms	14.28%
Total number of farms tested for H5	72
Number of Positive AIV-H5 farms	58
Percentage of Positive AIV-H5 farms	80.55%
Number of negative AIV-H5 farms	14
Percentage of negative AIV-H5 farms	19.45%

Tracheal gross lesions

The main gross lesions in trachea of infected birds were severe congestion and hemorrhage in comparison with normal trachea (Figure 2).

Real-time Reverse Transcriptase/Polymerase Chain Reaction RRT-PCR

All positive samples with AIV-type A and then AIV-H5 were also positive for RRT-PCR in comparison with +ve and -ve control (Figure 3).

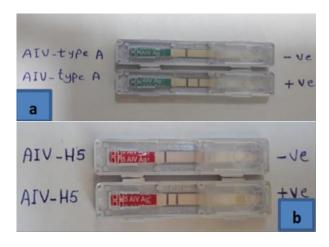


Figure 1: Positive and negative samples for AIV-A and H5 antigen using rapid chromatographic strip.

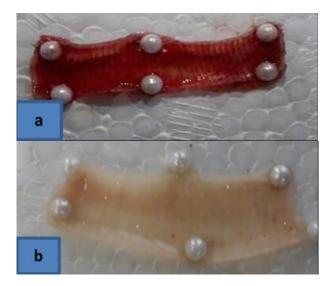


Figure 2: (a) Congestion and hemorrhage of infected trachea, (b) normal trachea.

Histopathological finding Trachea

The histopathological examination of infected trachea specimens showed obvious desquamation of lining epithelium with complete loss of cilia, extensive hemorrhage and infiltration of inflammatory cells especially macrophages, lymphocytes and plasma cells associated with congestion of blood vessels in lamina propria (Figure 4b) compared with control trachea in which the cilia are still intact (Figure 4a). Degeneration and necrosis of chondrocytes was evident in the hyaline cartilage (Figure 4c). Infiltration of inflammatory cells with coagulative necrosis and edema was prominent in the muscular layer (Figure 4d).

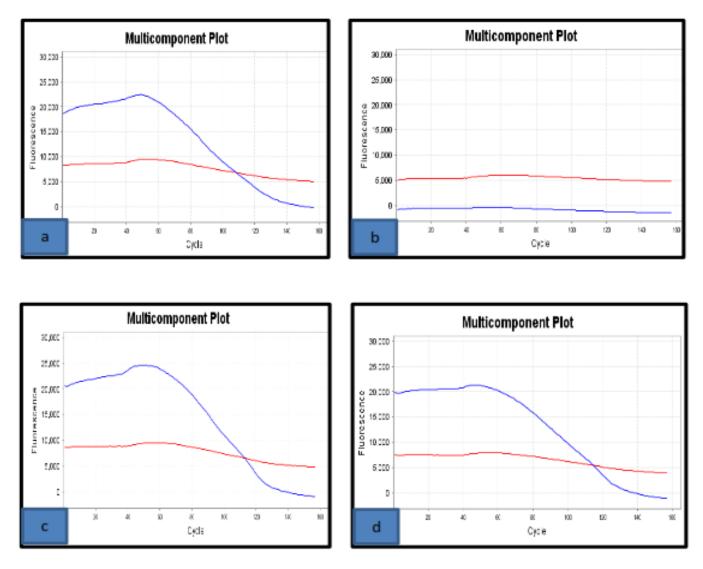


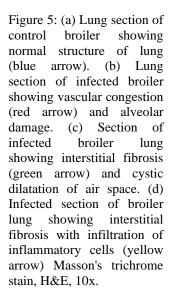
Figure 3: RRT-PCR assay examining cDNA from different samples infected with AIV-H5, a = +ve control, b = -ve control, c and d = 2 out of 8 +ve samples.

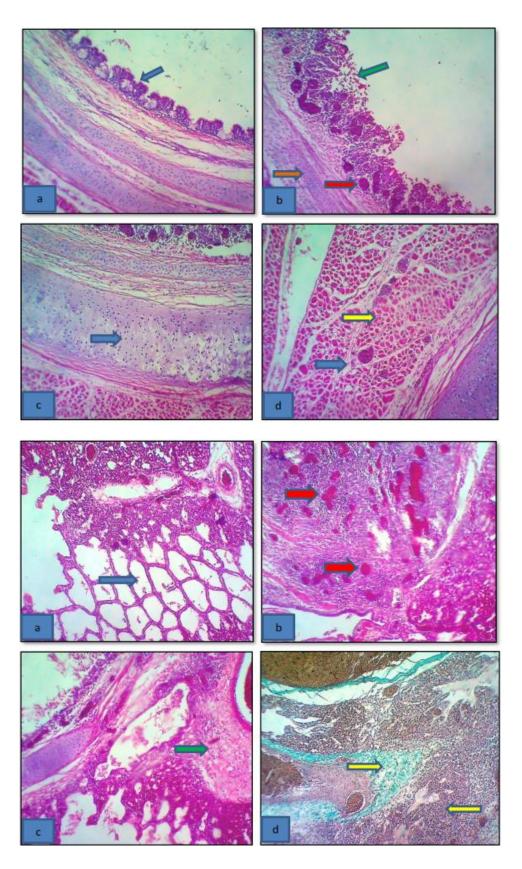
Lung

Histopathological sections of infected lungs revealed diffuse alveolar damage and severe multifocal vascular congestion with extensive hemorrhage (Figure 5b). Interstitial fibrosis leads to thickening of interlobular septa, cystic dilatation of the air space and edema (Figure 5c). Interstitial fibrosis with infiltration of lymphocytes, macrophages and heterophils in the interstitial and alveolar lumen (Figure 5d). Normal section of lung (Figure 5a).

Spleen

The histopathological changes of spleen from infected birds is characterized by multifocal necrosis of lymphoid cells with thickening of blood vessels wall lead to lumen stenosis (Figure 6a), was a deposition of fibrinous material in the splenic tissue associated with lymphocytes depletion and disappearance of the germinal centers (Figure 6b, c). Figure 4: (a) Tracheal section of healthy broiler showing normal structure of trachea (blue arrow). (b) Tracheal section of infected broiler with AIV-H5 virus showing desquamation of epithelium lining with complete loss of cilia (green arrow) hemorrhage (red infiltration arrow) of inflammatory cells (organ arrow) and congestion of blood vessels. (c) Tracheal section of infected broiler with AIV-H5 virus showing degeneration (blue arrow). (d) Necrosis of chondrocytes of hyaline cartilage (blue arrow) coagulative necrosis and edema of muscular layer (yellow arrow). H&E, 10x.





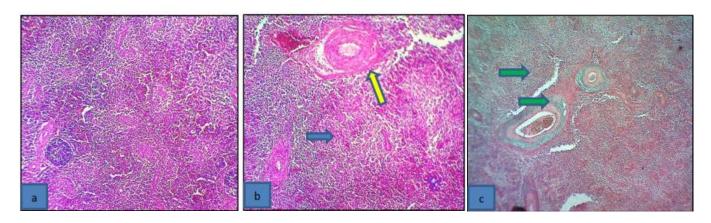


Figure 6: (a) Spleen section of control broiler showing normal architecture. (b) Spleen section of infected broiler showing focal area of necrosis of lymphoid cells (blue arrow) thickening of blood vessels wall and deposition of fibrinous material (yellow arrow), H&E, 10x. (c) Spleen section of infected broiler showing deposition of fibrinous material within splenic tissue and around the blood vessels walls (green arrow). Masson's trichrome stain, H&E, 4x.

Discussion

During last three decades, there has been a noticeable increase in the occurring of highly pathogenic avian influenza virus (HPAI) outbreaks and in the numbers of avian species infected in those outbreaks (11). Viruses of subtypes H5 and H7 which classified as highly pathogenic avian influenza (HPAI) characteristically produce fatal and severe systemic infection with high mortality rate in poultry farms. The HPAI viruses replicates within the endothelial cells of target organs throughout the vascular system of chicken during primary infection (12). The pathogenesis of infection include invasion of virus to the blood-brain barrier which followed by infection of nervous system with concurrently attacking and replication in different parenchymal tissues and organs (13). The RRT-PCR assay has been used worldwide successfully for the detection of many pathogens including RNA viruses (5). Since the rapid diagnostic test kits which are used for the detection of the nucleoprotein of the virus do not identify the subtypes of influenza A virus. The percentage of infected farms with this type of test was 85.72%, However, in the present research, the rapid chromatographic kit was used to detect the HA spikes of H5 antigen gave a percentage of 80.55% in tested farms, this study agreed with Chua et al (14) who mentioned ranges from 34.4% to 81.3%. Moreover, the rapid feature of these diagnostic kits could accomplish within 10-15 min and need no specific indicators, with high sensitivity and specificity and its significance for urgent control measures of H5 HPAI viruses (15). In microscopical examination the severe hemorrhage of trachea agreed with Kim et al (16) and the congestion occur as a result of excessing the capillary vessels with infiltration of leukocytes in the lamina properia (17). In comparison to the previous study Spackman et al (18), the methods which

are used in the present study provide many benefits over classical diagnostic methods, including rapid, easy, accurate and flexible. The RRT-PCR assay provides several benefits than standard assay, since it is rapid test for detection of the AIVs subtypes, including identification of co-circulating viruses (19). The one-step RRT-PCR assay is sensitive and it did not need to isolate and cultivate of pathogens from clinical samples. In conclusion the RRT-PCR assay described here provides a quick, sensitive and specific method to detect the H5 gene of AIVs in the clinical samples of diseased poultry (20). The pathological changes include anatomical and histological changes which occur as a result of viral invasion and replication within the target cells of susceptible host tissue or organ (21). The replication of AIV-H5 within the respiratory epithelial cells leads to development and occurrence of histopathological changes of trachea and lung. In the previous study which showed that the AIV carried by vascular endothelial cells of infected birds lead to alteration in the permeability of blood vessels resulting in hemorrhage, infiltration of inflammatory cells and necrotic lesions in lung, trachea and spleen (22,23).

Conclusion

In conclusion AIV-H5 infection cause severe pathological and histopathological changes as a result of systemic infection, the rapid chromatographic strips showed high specificity and sensitivity as field diagnostic tool. The RRT-PCR assay was highly sensitive and specific for detection of avian influenza virus subtypes with highly pathogenic properties (HPAI).

Acknowledgement

Authors would like to thanks the College of Veterinary medicine .University of Mosul / Iraq.

References

- Chen H, Smith GJ, Zhang SY, Qin K, Wang J, Li KS, Webster RG, Peiris JS, Guan Y. Avian flu: H5N1 virus outbreak in migratory waterfowl. Nature. 2005;436:191-192. doi:10.1038/nature03974
- Group WOFHNEW. Toward a unified nomenclature system for highly pathogenic avian influenza virus (H5N1). Emerg Infect Dis. 2008;14(7):e1. doi: 10.3201/eid1407.071681.
- Cagle C, Wasilenko J, Adams SC, Cardona CJ, To TL, Nguyen T, Spackman E, Suarez DL, Smith D, Shepherd E, Roth J, Pantin MJ. Differences in pathogenicity, response to vaccination, and innate immune responses in different types of ducks infected with a virulent H5N1 highly pathogenic avian influenza virus from Vietnam. Avian Dis. 2012;56(3):479-87. doi: 10.1637/10030-120511
- Alexander DJ. A review of avian influenza in different bird species. Vet Microbiol. 2000;74(1):3-13. Doi: 10.1016/s0378-1135(00)00160-7.
- WHO. Recommended laboratory tests to identify avian influenza A virus in specimens from humans. Geneva .June. 2005; http://www.who.int/csr/disease/avian_influenza/guidelines/avian_labt ests2.
- Chen W, Xu Z, Mu J, He B, Yang L, Lin L, Meng S, Mu F, Gan H. Real-time quantitative fluorescent reverse transcriptase-PCR for detection of severe acute respiratory syndrome associated coronavirus RNA. Mol Diagn. 2004;8:231-235. doi: 10.1007/bf03260067
- Rabee R.H.S. and Jaber I.M. The interference of Newcastle, avian influenza and infectious bursal disease vaccines with the efficiency of IB vaccine in broiler chicks by using quantitative RT-PCR test. Iraqi Journal of Veterinary Sciences.2019.33(1):105-109.
- Rashid S, Naeem K, Ahmed Z, Saddique N, Abbas MA, Malik SA. Multiplex polymerase chain reaction for the detection and differentiation of avian influenza viruses and other poultry respiratory pathogens. Poult Sci. 2009;88:2526-2531. doi: 10.3382/ps.2009-00262
- Luna LG. Manual of histologic staining methods of the armed forces institute of pathology. 3rd ed. New York: McGraw-Hill; 1968.38-76.
- Al-Bajaril SH A, Al- Akash1 M A, Ismail H KH. Experimental detection of antioxidant and atherogenic effects of grapes seeds extracts in rabbits. Iraqi Journal of Veterinary Sciences.2019.33(2):243-249.
- Kapczynski DR, Pantin M, Guzman SG, Ricardez Y, Spackman E, Bertran K. Characterization of the 2012 highly pathogenic avian influenza H7N3 virus isolated from poultry in an outbreak in Mexico: pathobiology and vaccine protection. J Virol. 2013;87(16):9086-96. doi: 10.1128/JVI.00666-13

- Brown CC, Olander HJ, Senne DA. A pathogenesis study of highly pathogenic avian influenza H5N2 in chickens, using immunohistochemistry. J Comp Pathol. 1992;107:341-348. doi.org/10.1354/vp.44-5-635
- Silvano FD, Yoshikawa M, Shimada A, Otsuki K, Umemura T. Enhanced neuro- pathogenicity of avian influenza A virus by passages through air sac and brain of chicks. J Vet Med Sci. 1997;59:143-148. doi.org/10.1292/jvms.59.143
- Chua TH, Ellis TM, Wong CW, Guan Y, Ge, SX, Peng G, Lamichane C, Maliadis C, Tan SW, Selleck P, Parkinson J. Performance evaluation of five detection test for avian influenza antigen with various avian sample. Avian Dis. 2007;56: 96105. doi:10.1637/0005-2086(2007)051[0096:PEOFDT]2.0.CO;2
- Nguyen LT, Nakaishi K, Motojima K, Ohkawara A, Minato F, Maruyama J, Hiono T, Matsuno K, Okamatsu M, Kimura T, Takada A, Kida H, Sakoda Y. Rapid and broad detection of H5 hemagglutinin by an immune chromatographic kit using novel monoclonal antibody against highly pathogenic avian influenza virus belonging to the genetic clade. PLoS One. 2017;12(8):e0182228. doi: 10.1371/journal.pone.0182228
- Kim HR, Kwon YK, Jang I, Lee YJ, Kang HM, Lee EK. Pathologic changes in wild birds infected with highly pathogenic avian influenza A (H5N8) Viruses, South Korea, 2014. Emerg Infect Dis. 2015;21(5):775-80. doi: 10.3201/eid2105.141967
- Ali A, Elmowalid G, Abdel-Glil1M, Sharafeldin A, Abdallah F, Mansour S, Nagy A, Ahmed B, Abdelmoneim M. Etiology and pathology of epidemic outbreaks of avian influenza H5N1 infection in Egyptian chicken farms. Polish J Vet Sci. 2015;18(4):779-786. doi: 10.1515/pjvs-2015-0101
- Spackman EDA, Senne TJ, Myers LL, Bulaga LP, Garber ML, Perdue K, Lohman LT, Daum DL. Development of a real-time reverse transcriptase PCR assay for type A influenza virus and the avian H5 and H7 hemagglutinin subtypes. J Clin Microbiol. 2002;40:3256-3260. doi: 10.1637/0005-2086-47.s3.1079
- Isabella M, Silvia O, Annalisa S, Cristian DB, Francesca B, Angela S, Alessandra D, Bianca Z, Ilaria C, Giovanni C. Development and validation of a one-step Real-Time PCR assay for simultaneous detection of subtype H5, H7, and H9 avian influenza viruses. J Clin Micro. 2008;1769-177. doi: 10.1128/JCM.02204-07
- Tan ML, Niu Y, Shui W, Lin J, Li M Zhang C. The establishment of real-time fluorescent quantitative polymerase chain reaction (PCR) for detection of highly pathogenic avian influenza virus subtype H5N1. J Immunol Tech Infect Dis. 2017;6(1):1. doi: 10.4172/2329-9541.1000154
- Swayne DE. Understanding the complex pathobiology of high pathogenicity avian influenza viruses in birds. Avi Dis. 2007;51(1 Suppl):242-9. doi: 10.1637/7763-110706-REGR.1
- Pantin MJ, Swayne DE. Pathobiology of asian highly pathogenic avian influenza H5N1 virus infections in ducks. Avi Dis. 2007;51(1 Suppl):250-9. doi: 10.1637/7710-090606R.1
- Kuiken T, van den Brand J, van Riel D, Pantin M, Swayne DE. Comparative pathology of select agent influenza a virus infection. Vet Pathol. 2010;47(5):893-914. doi: 10.1177/0300985810378651.