

## Case Report

### Isolation and identification of HPAI from fertile egg derived from broiler breeder farm after natural vaccination breakage during 2010 in Egypt.

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#### Abstract

In May 2010, during routine monitoring of a commercial broiler breeder flock consists of 6 horizontal houses with total capacity 125000 bird. The farm were vaccinated 3 times with an inactivated H5N1 vaccine. The faerm added under weekly observation and monitoring for continuous 5 weeks after isolation and identification HPAI H5N1 virus from cloacal and oropharyngeal swabs. Different 10 cloacal, oropharyngeal swabs and fertile eggs “after separation of shell, albumin and yolk” samples were directed for AI common gene detection using RTq-PCR test and virus isolation using 10 days old SPF ECE in pooled samples (5 samples/ pool). Moreover 20 blood samples were collected for H5 antibody detection using HI test.

AI H5N1 virus were detected and isolated from different egg parts (egg shell, egg albumin and egg yolk) from the different 6 houses since 2<sup>nd</sup> weeks post natural infection and stopped after 7 weeks post infection in case of house 5 and 6. The mortality rate was range (27 to 43%) and the drops in egg quality and egg production (20 to 41%) were also recorded. In addition, the HI titers were 3.5 and 5.6 log<sub>2</sub> during first 2 weeks of infection and reach to 10log<sub>2</sub> at 4<sup>th</sup> week post infection. The isolated H5N1 virus from fertile eggs showed high nucleotides and amino-acid identities to recently variant HPAI H5N1 subclade 2.2.1 viruses. The vaccinal break seems to be associated with immune escape mutants and/or improper vaccination. The role of contaminated eggs as a source of infection and as a vehicle for spread of the virus should be considered in area with avian influenza outbreaks, possibilities of vertical transmission weren't studied during this study.

**Key words:** “Vaccine, “HPAI” Highly Pathogenic Avian Influenza, “AI” Avian Influenza, Egypt, broiler breeder and fertile egg”.

#### Introduction

Highly pathogenic avian influenza (HPAI) H5N1 virus infection is a devastating viral disease causing severe losses in the poultry industry worldwide. AIV are segmented negative-sense RNA viruses belong to the family Orthomyxoviridae that are divided into five genera, including influenza types A, B, and C, Isavirus and Thogotovirus **Palese and Shaw, (2007)**. Based on the hemagglutinin (HA) and

neuraminidase (NA), two main surface transmembrane glycoproteins, there are at least 17 HA (denoted H1 to H17) and probably 10 NA subtypes (denoted N1–N10) **Tonga et al. (2012)**.

The main control measures in the face of an AI outbreak are continuous massive surveillance, proper hygienic culling of infected birds, restriction of movement, and enforcement of biosecurity **Tian et al. (2005) and Capua and**

**Alexander, (2006).** But in case of developing countries, implementation of massive vaccination policy using H5-based inactivated vaccines as supportive control tool was implemented in several countries. Since the incursion of HPAI H5N1 in Egypt in mid-February 2006 **Aly *et al.* (2007); Kilany, (2007) and Safwat, (2007)**, Egypt have been embarked on mass uncontrolled vaccination program for commercial poultry with different types of inactivated H5N1 and H5N2 vaccines with wide variation in standards was implemented as the main control practice **Abdelwhab and Hafez, (2011)**. However, since 2007, infection of vaccinated flocks has increased and subsequent depopulation of positive flocks has not been uncommon **Aly *et al.* (2008)**. By late 2007, H5N1 outbreaks in vaccinated poultry in Egypt have been dramatically increased which criticized the effectiveness of the implemented vaccines **Peyre *et al.* (2009)**. Genetic and antigenic analysis of the circulating field viruses in vaccinated flocks revealed subtle antigenic variation and antigenic drift of the virus in comparison to vaccine strains **Arafa *et al.* (2010a,b); Hafez *et al.* (2010); Abdel-Moneim *et al.* (2011a,b); Abdelwhab *et al.* (2011); Grund *et al.* (2011); Abdelwhab *et al.* (2012)**. These mutations allowed the virus to evade the repertoire of host immune response after vaccination.

It was thought that due to the short clinical course of HPAI infected chickens as they usually die within 1–3 days, so it is unlikely to disseminate the HPAI virus in produced eggs via infected chickens and couldn't be detected. But a lot of studies were able to detect AIV in eggs of naturally as well as experimentally infected birds have been previously described **Bean *et al.* (1985); Cappucci *et al.* (1985); Kilany *et al.* (2011) and Abdelwhab *et al.* (2012)**.

Also in this study, we studied the possibilities to isolate HPAI H5N1 virus from fertile eggs obtained from naturally infected vaccinated broiler breeder chickens in Egypt during May 2010.

## Materials and Methods.

### Case report .

During May 2010, Broiler breeder flock consists of 125,000 divided on 6 different poultry houses, with same breed "Hubbard" and same age 32 week, was suffering from HPAI two weeks before starting the sampling due to late reporting. The farm was vaccinated 3 times with H5N1 vaccine seeded by A/Goose/Guangdong/1/1996/H5N1 (Re-1, Harbin, China) as following (1<sup>st</sup> vaccination at 10 days old, 2<sup>nd</sup> after one month and 3<sup>rd</sup> vaccination at 13 weeks old). The diseased birds were showed cyanosis comb and wattles as well as hemorrhages on the shank, facial edema and nervous manifestation at late stage of infection since 5<sup>th</sup> week post infection. The drop in egg production in case of HPAI is transient and was ranged from 20 to 41%, and an increase of the number of misshaped eggs (depigmentations, soft eggshell, shell-less and rough eggs). At the same time, mortality percent range from 27 to 43% (**Chart 2**). Samples for laboratory investigation were from each house, 10 tracheal swabs, 10 cloacal swabs, and 10 shelled eggs were collected randomly at weekly basis 14 days after the onset of clinical signs. At the same time, 20 serum samples were collected from each house. For further laboratory investigation, 5 swabs were pooled according to **OIE, (2009)**. From the collected eggs, eggshell wash, albumin, and yolk materials were separated, and each part was tested in 5 pool protocol.

### Virus isolation and detection.

Virus isolation carried out using 10-day-old specific-pathogen-free (SPF) hatching eggs via allantoic sac inoculation, **OIE, (2009)**. The HPAI H5N1 virus was isolated after first passage from tracheal and cloacal swabs collected from each house on weekly basis and confirmed by real-time reverse transcription–polymerase chain reaction (RT-qPCR). the HPAI H5N1 was recorded in house 5 and house 6 since 1<sup>st</sup> week of examination then the virus have been recorded in house 3, house 4, and in case of house 2 and house 1 detection of AI

was after 3<sup>rd</sup> week of examination. The AIV disease remain in the farm for more than 7 weeks post natural infection, also we record presence of H5 virus in different egg parts including egg shell, egg albumin and egg yolk in different houses and the dissemination of AIV virus in egg content start from 2<sup>nd</sup> week post natural infection and stops after 5<sup>th</sup> weeks post infection in case of house 5 and 6 (**Table1, 2 and 3**).

#### **RT-qPCR for H5N1 VIRAL RNA detection.**

RNA was extracted from 5 sample pools using a MagNA Pure LC Total Nucleic Acid Extraction kit following manufacturer's instructions and a MagNA Pure LC instrument (Roche, Mannheim, Germany). The matrix gene of avian influenza type A viruses was amplified from RNA samples using a one-step Real-Time PCR Kit (Qiagen, Valencia, CA) as described by (**Spackman et al., 2002**), and the test was conducted in a Stratagene MX3005P real-time PCR machine (Stratagene, Agilent Technologies, Santa Clara, CA). Detection of H5N1 subtype was done by avian influenza virus H5N1 Real-Time RT-PCR Kit (Roche, Mannheim, Germany), according to the manufacturer's instructions, using the Light Cycler H 2.0 machine (Roche, Mannheim, Germany). The H5N1 viral RNA could be detected in cloacal and tracheal swabs as well as from eggshell, egg albumin and yolk materials collected from affected houses.

**Serological investigation.** The detection of avian influenza H5- specific antibodies in the serum samples using hemagglutination inhibition (HI) test was performed according to **OIE, (2009)** using 4 hemagglutinin units of homologous A/Goose/ Guandong/1/1996 H5N1 antigen, supplied by the vaccine-producing company. Serologic examination showed the HI titers ranging from 3.5 to 5.6 log<sub>2</sub> during 1<sup>st</sup> week of monitoring in house 6 and house 5. On the other hand, on other houses 3, 4, 2 and 1 was not tested at first week; moreover HI titers reach to 10 log<sub>2</sub> in all houses after 3<sup>rd</sup> week of examination (**Table 2**).

#### **Sequence and phylogentic analysis.**

Sequencing of the H5 gene of the virus isolated

from cloacal, oropharyngeal swabs and designated as A/chicken/Egypt/10127s-NLQP/2010 (H5N1) was conducted as previously mentioned by **Arafa et al. (2010a)** using BigDye Terminator v3.1 Cycle Sequencing Kit on an automatic sequencer (ABI-3130; Applied Biosystems, Foster City, CA). Sequence data were analyzed and edited with BioEdit version 7.0.9.0 **Hall, (1999)**. Amino-acid sequences were deduced and a BLASTN search was performed to identify the query sequence and to find similar sequences. The H5 gene sequence of the isolated virus in this study was submitted to GenBank, and the accession number is HQ198274. Phylogenetic analyses for the full length of the H5 genes and the highest homologous H5 gene sequences obtained after blast were produced based on CLUSTAL V multiple sequence alignment program, version 1.83 of MegAlign module of Lasergene DNASTar software (Madison, WI) to determine nucleotide and amino acid sequence similarities and relationships, with 100 bootstrap using a GTR+G model as recommended by ModelTest (Topali). The tree was drawn with Dendroscope **Huson et al. (2007)**.

The virus had polybasic cleavage motif PQGERRRKKRGLF, which indicates high pathogenic subtype as for clade 2.2.1 viruses. The isolated virus showed 99.8% and 99.7% nucleotides and amino-acid identities, respectively, and clustered mainly with viruses from recently confirmed H5N1 variant viruses and with currently circulating viruses in commercial birds and vaccinated commercial farms from different places in Egypt in 2010.

**Table 1.** Results of laboratory investigation in house 1 and 2.

| Samples type                                            | hs1 |      |      |       |      | hs2  |      |      |      |      |
|---------------------------------------------------------|-----|------|------|-------|------|------|------|------|------|------|
|                                                         | W1  | W2   | W3   | W4    | W5   | W1   | W2   | W3   | W4   | W5   |
| <i>Virus isolation</i>                                  |     |      |      |       |      |      |      |      |      |      |
| Tracheal swab                                           | NT  | Neg  | Neg  | Post  | Post | NT   | Neg  | Neg  | Post | Post |
| Cloacal swab                                            | NT  | Neg  | Neg  | Post  | Post | NT   | Neg  | Neg  | Post | Post |
| Egg shell                                               | NT  | Neg  | Neg  | Post  | Post | NT   | Neg  | Neg  | Post | Post |
| Egg albumin                                             | NT  | Neg  | Neg  | Post  | Post | NT   | Neg  | Neg  | Post | Neg  |
| Egg yolk                                                | NT  | Neg  | Neg  | Neg   | Neg  | NT   | Neg  | Neg  | Neg  | Post |
| <i>RT-Qpcr</i>                                          |     |      |      |       |      |      |      |      |      |      |
| Tracheal swab                                           | NT  | Neg  | Neg  | Post  | Post | NT   | Neg  | Neg  | Post | Post |
| Cloacal swab                                            | NT  | Neg  | Neg  | Post  | Post | NT   | Neg  | Neg  | Post | Post |
| Egg shell                                               | NT  | Neg  | Neg  | Post  | Post | NT   | Neg  | Neg  | Post | Post |
| Egg albumin                                             | NT  | Neg  | Neg  | Post  | Post | NT   | Neg  | Neg  | Post | Neg  |
| Egg yolk                                                | NT  | Neg  | Neg  | Neg   | Neg  | NT   | Neg  | Neg  | Neg  | Post |
| <i>Serology HI test</i>                                 |     |      |      |       |      |      |      |      |      |      |
| HI mean titer                                           | NT  | 8.5  | 9    | 10    | 10   | NT   | 8.3  | 9.5  | 10   | 10   |
| STD $\pm$                                               | NT  | 0.46 | 0.3  | 0.24  | 0.2  | NT   | 0.9  | 0.76 | 0.32 | 0.4  |
| Post %                                                  | NT  | 100% | 100% | 100%  | 100% | NT   | 100% | 100% | 100% | 100% |
| Mortality                                               | 2%  | 2.6% | 10%  | 15.8% | 23%  | 1.2% | 4%   | 11%  | 21%  | 27%  |
| <i>Egg production in comparison with standard curve</i> |     |      |      |       |      |      |      |      |      |      |
|                                                         | 88% | 89%  | 70%  | 65%   | 54%  | 89%  | 92%  | 74%  | 53%  | 56%  |

hs= poultry house

NT= Not tested

**Table 2.** Results of laboratory investigation in house 3 and 4.

| Samples type                                     | hs3 |      |      |       |      | hs4  |      |      |      |      |
|--------------------------------------------------|-----|------|------|-------|------|------|------|------|------|------|
|                                                  | W1  | W2   | W3   | W4    | W5   | W1   | W2   | W3   | W4   | W5   |
| Virus isolation                                  |     |      |      |       |      |      |      |      |      |      |
| Tracheal swab                                    | NT  | Post | Post | Post  | Post | NT   | Post | Post | Post | Post |
| Cloacal swab                                     | NT  | Post | Post | Post  | Post | NT   | Post | Post | Post | Post |
| Egg shell                                        | NT  | Post | Post | Post  | Post | NT   | Post | Post | Post | Post |
| Egg albumin                                      | NT  | Post | Post | Post  | Neg  | NT   | Neg  | Post | Post | Neg  |
| Egg yolk                                         | NT  | Neg  | Post | Post  | Neg  | NT   | Neg  | Post | Post | Post |
| RRT-PCR                                          |     |      |      |       |      |      |      |      |      |      |
| Tracheal swab                                    | NT  | Post | Post | Post  | Post | NT   | Post | Post | Post | Post |
| Cloacal swab                                     | NT  | Post | Post | Post  | Post | NT   | Post | Post | Post | Post |
| Egg shell                                        | NT  | Post | Post | Post  | Post | NT   | Post | Post | Post | Post |
| Egg albumin                                      | NT  | Post | Post | Post  | Neg  | NT   | Neg  | Post | Post | Neg  |
| Egg yolk                                         | NT  | Neg  | Post | Post  | Neg  | NT   | Neg  | Post | Post | Post |
| Serology HI test                                 |     |      |      |       |      |      |      |      |      |      |
| HI mean titer                                    | NT  | 6.5  | 9.3  | 10    | 10   | NT   | 7.6  | 9.2  | 10   | 10   |
| STD $\pm$                                        | NT  | 0.35 | 0.87 | 0.12  | 0.1  | NT   | 1.2  | 1.3  | 0.2  | 0.1  |
| Post %                                           | NT  | 100% | 100% | 100%  | 100% | NT   | 100% | 100% | 100% | 100% |
| Mortality                                        | 8%  | 9.6% | 13%  | 25.8% | 34%  | 6.3% | 9.9% | 15%  | 23%  | 34%  |
| Egg production in comparison with standard curve |     |      |      |       |      |      |      |      |      |      |
|                                                  | 75% | 75%  | 65%  | 64%   | 53%  | 75%  | 73%  | 64%  | 53%  | 59%  |

hs= poultry house

NT= Not tested

**Table 3.** Results of laboratory investigation In house 5 and 6.

| Samples type                                     | hs5  |       |      |      |      | hs6  |       |       |      |      |
|--------------------------------------------------|------|-------|------|------|------|------|-------|-------|------|------|
|                                                  | W1   | W2    | W3   | W4   | W5   | W1   | W2    | W3    | W4   | W5   |
| Virus isolation                                  |      |       |      |      |      |      |       |       |      |      |
| Tracheal swab                                    | Post | Post  | Post | Post | Neg  | Post | Post  | Post  | Post | Neg  |
| Cloacal swab                                     | Post | Post  | Post | Post | Neg  | Post | Post  | Post  | Post | Neg  |
| Egg shell                                        | Post | Post  | Post | Post | Neg  | Post | Post  | Post  | Post | Neg  |
| Egg albumin                                      | Post | Post  | Post | Post | Neg  | Neg  | Neg   | Post  | Post | Neg  |
| Egg yolk                                         | Neg  | Neg   | Post | Post | Neg  | Neg  | Neg   | Post  | Post | Neg  |
| RRT-PCR                                          |      |       |      |      |      |      |       |       |      |      |
| Tracheal swab                                    | Post | Post  | Post | Post | Neg  | Post | Post  | Post  | Post | Neg  |
| Cloacal swab                                     | Post | Post  | Post | Post | Neg  | Post | Post  | Post  | Post | Neg  |
| Egg shell                                        | Post | Post  | Post | Post | Neg  | Post | Post  | Post  | Post | Neg  |
| Egg albumin                                      | Post | Post  | Post | Post | Neg  | Neg  | Neg   | Post  | Post | Neg  |
| Egg yolk                                         | Neg  | Neg   | Post | Post | Neg  | Neg  | Neg   | Post  | Post | Neg  |
| Serology HI test                                 |      |       |      |      |      |      |       |       |      |      |
| HI mean titer                                    | 5.6  | 6.5   | 10   | 10   | 10   | 3.5  | 7.6   | 10    | 10   | 10   |
| STD $\pm$                                        | 2.2  | 1.35  | 0.87 | 0    | 0    | 2.6  | 2.2   | 0     | 0    | 0    |
| Post %                                           | 80%  | 100%  | 100% | 100% | 100% | 60%  | 100%  | 100%  | 100% | 100% |
| Mortality                                        | 17%  | 23.5% | 29%  | 37%  | 38%  | 18%  | 22.2% | 27.9% | 38%  | 43%  |
| Egg production in comparison with standard curve |      |       |      |      |      |      |       |       |      |      |
|                                                  | 65%  | 53%   | 65%  | 68%  | 75%  | 55%  | 59%   | 64%   | 74%  | 80%  |

hs= poultry house

NT= Not tested

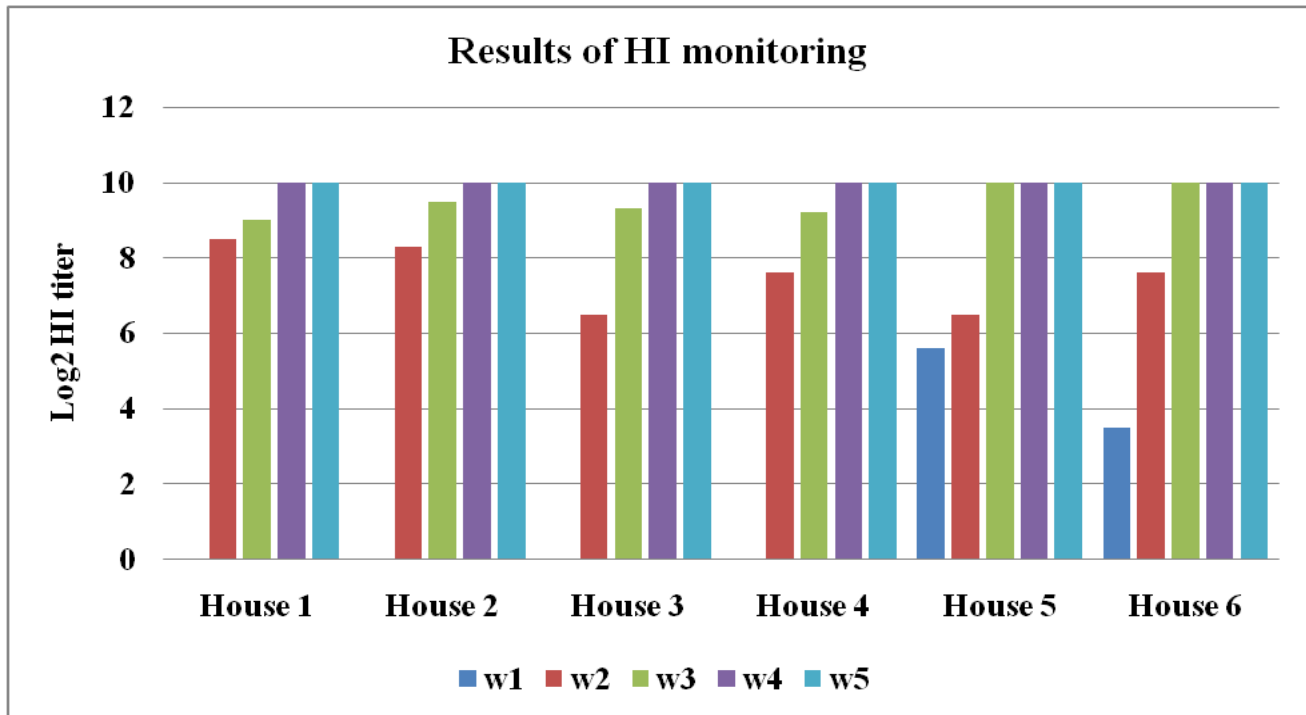


Chart 1. Results of HI testing during the 5 tested weeks.

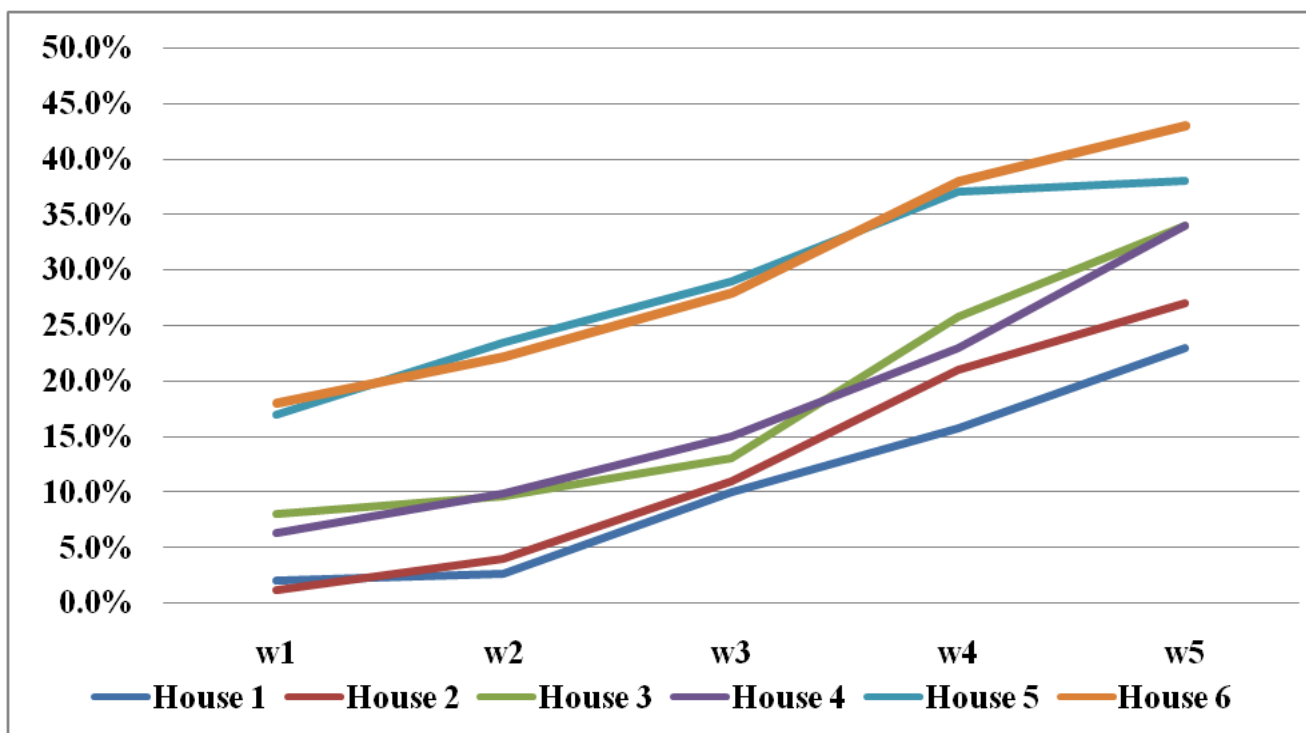
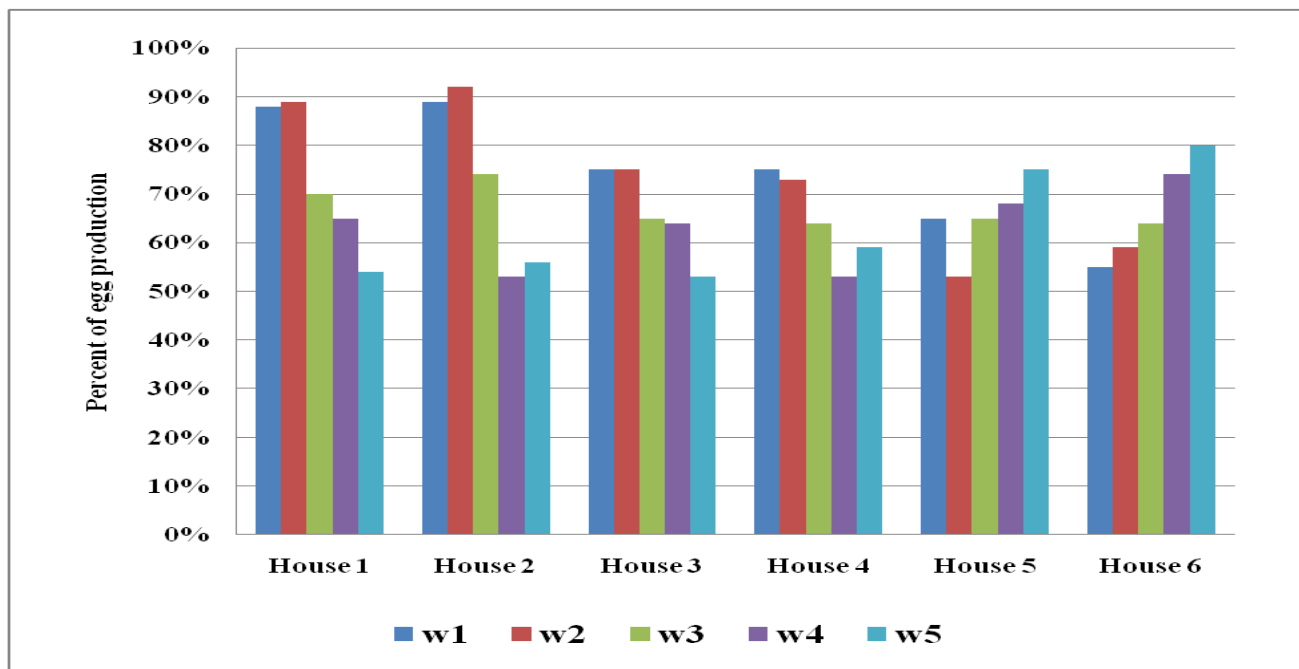


Chart 2. Cumulative Mortality.



**Chart 3.** Drop in egg production curve during 5 weeks of examination

### Discussion.

HPAI H5N1 outbreaks in vaccinated poultry flocks have been reported in many countries and there were many factors are incriminated as a cause of the vaccinal breaks, such as incomplete immunization coverage and/or vaccine failure, as described by **Simth *et al.* (2006)** in China, **Aly *et al.* (2008)** in Egypt and **Henning *et al.* (2009)** in Vietnam. The Continuous emerging of the endemic HPAI H5N1 virus within vaccinated birds in Egypt in the last five years and emerging of different antigenic and genetic mutant viruses as in group B (B1, B2.1 and B2.2) **Kilany, (2013)**. So far, two different viruses are circulating now in Egypt; subgroup A2 “classic 2.2.1” strains infect mostly backyard birds and human and the newly emerging subgroup B2.2 “2.2.1 variants” circulate mainly in vaccinated commercial farms in Egypt, circulation of genetically and antigenically distinct H5N1 viruses in vaccinated poultry flocks is common **Abdelwhab *et al.* (2010)**; **Arafa *et al.* (2010 a, b)**; **Balish *et al.* (2010)**, **Abdelwhab *et al.* (2011)** and

### **Kilany *et al.* (2011).**

The virus RNA were detected from eggshell, albumin and egg yolk by RT-q PCR, and the isolation of the virus was successful from egg shell, egg albumin and yolk as shown in (Table 1, 2 and 3) which is in accordance with recently reported results **Pillai, *et al.* (2010)** from experimentally infected turkey by H3N2 AIV, moreover **Bean, *et al.* (1985)**; **Cappucci *et al.* (1985)**; **Li *et al.* (2006)**; **Kilany *et al.* (2010)**; **Abdelwhab *et al.* (2012)** isolate AIV from internal contents of eggs obtained from naturally or experimentally infected birds. Also, HPAI H5N1 virus was recovered from the yolk and albumin of naturally infected Japanese quail **Promkuntod *et al.* (2006)** and egg shells of ducks and geese **Li *et al.* (2006)**.

The role of contaminated eggs as a source of infection and as a vehicle for spread of the virus should be considered in area with avian influenza outbreaks **Kilany *et al.* (2010)**; **Pillai *et al.* (2010)**; **Abdelwhab *et al.* (2011)**. However proof of vertical transmission is lacking. Moreover, HPAIV is usually embryo lethal and

hatching of internally contaminated eggs is unlikely **Easterday *et al.* (1997)**.

The mortality rate during this study reach (27 to 43%) and egg production and egg quality dropped (20 to 41%). On the other hand, serologic examination of sera collected two weeks after the onset of the clinical signs, the HI titers in the first affected two houses (House 6 and 5) were 5.6 and 3.5 log<sub>2</sub>, which indicate improper vaccination **Tian *et al.* (2005)** and the HI titers shows repaid poster since week 3 post infection in all houses and reach to 10 log<sub>2</sub>, using the homologous H5N1 antigen.

The isolated virus had polybasic cleavage motif ERRRKRR/GLF, which indicates high pathogenic subtype as for clade 2.2.1 viruses **Abdelwhab, (2012); Kilany, (2013)**. It has relatively high similarities to nucleotides and amino acids and clustered with viruses from recently confirmed HPAI H5N1 commercial poultry cases in Egypt. Moreover, several amino-acid substitutions of viral H5 protein were observed at different positions not shown in this study. These amino-acid residues are located mainly in the most exposed part of the globular head of the hemagglutinin **Kilany, (2013)**. In Egypt, circulation of genetically and antigenically distinct H5N1 viruses in vaccinated poultry flocks is not uncommon **Arafa *et al.* (2010 a, b); Balish *et al.* (2010), Abdelwhab *et al.* (2011) and Kilany *et al.* (2011)**.

Accordingly, in case of an outbreak of HPAI H5N1, regular disinfection of contaminated eggshell alone will be insufficient to disclose the potential hazard for spread of the virus through infected, contaminated cracked eggs, and egg flats during their marketing or from one farm to another by vendors and vehicles. Therefore, we recommend that eggs from infected birds should be culled in the farm to prevent further spread of the virus into the environment. In conclusion, although currently no available data suggest that the vertical transmission of HPAI but handling of contaminated eggs could be potential hazards of spreading AI infection to the surrounding environment by movement of eggs and contaminated egg trays in areas of an outbreak must be emphasized.

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## References

- Abdel-Moneim, A. S., M. A. Afifi, and M. F. El-Kady (2011a)**. Genetic drift evolution under vaccination pressure among H5N1 Egyptian isolates. *Virology J.*, 8:283.
- Abdel-Moneim, A. S., G. M. Shehab, and A. A. Abu-Elsaad (2011b)**. Molecular evolution of the six internal genes of H5N1 equine influenza A virus. *Archives of Virology* 156: 1257-1262.
- Abdelwhab, E. M., and H. M. Hafez. (2011)**. An overview of the epidemic of highly pathogenic H5N1 avian influenza virus in Egypt: epidemiology and control challenges. *Epidemiology and Infection* 139:647-657.
- Abdelwhab, E. M., A. Selim, A. Arafa, S. Galal, W. H. Kilany, M. K. Hassan, M. M. Aly, and H. M. Hafez (2010)**. Circulation of avian influenza H5N1 in live bird markets in Egypt. *Avian Dis.* 54:911-914.
- Abdelwhab, E. M., C. Grund, M. M. Aly, M. Beer, T. C. Harder, and H. M. Hafez (2011)**. Multiple dose vaccination with heterologous H5N2 vaccine: immune response and protection against variant clade 2.2.1 highly pathogenic avian influenza H5N1 in broiler breeder chickens. *Vaccine* 29:6219-6225.
- Abdelwhab, E. M., C. Grund, M. M. Aly, M. Beer, T. C. Harder, and H. M. Hafez (2012)**. Influence of maternal immunity on vaccine efficacy and susceptibility of one day old chicks against Egyptian highly pathogenic avian influenza H5N1. *Veterinary microbiology* 155:13-20.
- Aly, M. M., Z. Kanawaty, A. Arafa, W. H. Kilany, and E. M. Abdelwhab (2007)**. One-year surveillance on avian influenza H5N1 in backyard poultry in Egypt. *Proceedings of the 4th International Symposium of Turkey Production, Berlin, Germany*:294-306.
- Aly, M. M., A. Arafa, and M. K. Hassan**

- (2008). Epidemiological findings of outbreaks of disease caused by highly pathogenic H5N1 avian influenza virus in poultry in Egypt during 2006. *Avian Dis.* 52: 269–277.
- Arafa, A., D. L. Suarez, M. K. Hassan, and M. M. Aly (2010a).** Phylogenetic analysis of HA and NA genes of HPAI-H5N1 Egyptian strains isolated from 2006 to 2008 indicates heterogeneity with multiple distinct sublineages. *Avian Dis.* 54:345–349.
- Arafa, A. S., A. A. Selim, M. K. Hassan, and M. M. Aly (2010b).** Genetic characterization of variant strains of highly pathogenic avian influenza H5N1 that escaped detection by real-time reverse transcriptase-PCR diagnostic tests. *Avian diseases* 54:673–676.
- Balish, A. L., T. Davis, M. D. Saad, N. El-Sayed, H. Esmat, J. A. Tjaden, K. C. Earhart, L. E. Ahmed, M. Abd El-Halem, A. Hakem, A. M. Ali, S. A. Nassif, E. A. El-Ebiary, M. Taha, M. M. Aly, A. Arafa, E. O'Neill, X. Xiyan, N. J. Cox, R. O. Donis, and A. I. Klimov (2010).** Antigenic and genetic diversity of highly pathogenic avian influenza A (H5N1) viruses isolated in Egypt. *Avian Dis.* 54: 329–334.
- Bean, W. J., Y. Kawaoka, J. M. Wood, J. E. Pearson, and R. G. Webster (1985).** Characterization of virulent and avirulent A/chicken/Pennsylvania/ 83 influenza A viruses: potential role of defective interfering RNAs in nature. *J. Virol.* 54:151–160.
- Capua, I., and D. J. Alexander (2006).** The challenge of avian influenza to the veterinary community. *Avian pathology : Journal of the W.V.P.A* 35:189-205.
- Cappucci, D. T. Jr, D. C. Johnson, M. Brugh, T. M. Smith, C. F. Jackson, J. E. Pearson, and D. A. Senne (1985).** Isolation of avian influenza virus (subtype H5N2) from chicken eggs during a natural outbreak. *Avian Dis.* 29: 1195–2000.
- Cattoli, G., A. Fusaro, I. Monne, F. Coven, T. Joannis, H. S. El-Hamid, A. A. Hussein, C. Cornelius, N. M. Amarín, M. Mancin, E. C. Holmes, and I. Capua (2011a).** Evidence for differing evolutionary dynamics of A/H5N1 viruses among countries applying or not applying avian influenza vaccination in poultry. *Vaccine* 29:9368-9375.
- Easterday, B. C., V. S. Hinshaw, and D. A. Halvorson (1997).** Influenza, p. 583-605. *Disease of poultry* In B. W. Calnek, H. J. Barnes, C. W. Beard, L. R. McDougald, and Y. M. Saif (ed.), *Diseases of Poultry*, 10th ed. Iowa State University Press, Ames, Iowa.
- Grund, C., S. M. Abdelwhab el, A. S. Arafa, M. Ziller, M. K. Hassan, M. M. Aly, H. M. Hafez, T. C. Harder, and M. Beer (2011).** Highly pathogenic avian influenza virus H5N1 from Egypt escapes vaccine-induced immunity but confers clinical protection against a heterologous clade 2.2.1 Egyptian isolate. *Vaccine* 29:5567-5573.
- Hafez, M. H., A. Arafa, E. M. Abdelwhab, A. Selim, S. G. Khoulosy, M. K. Hassan, and M. M. Aly (2010).** Avian influenza H5N1 virus infections in vaccinated commercial and backyard poultry in Egypt. *Poultry science* 89:1609-1613.
- Hall, T. A (1999).** BioEdit: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. *Nucleic Acids Symp.Ser.* 41:95–98.
- Henning, J., D. U. Pfeiffer, and L. T. Vu (2009).** Risk factors and characteristics of H5N1 highly pathogenic avian influenza (HPAI) post vaccination outbreaks. *Vet. Res.* 40:15–27.
- Huson, D. H., D. C. Richter, C. Rausch, T. Dezulian, M. Franz, and R. Rupp.** Dendroscope: an interactive viewer for large phylogenetic trees. *BMC Bioinform.* 8:460.
- Kilany, W. H. (2007).** Studies on avian influenza virus. Master Thesis. Zagazige University, Zagazige, Egypt.
- Kilany, W.H (2013)** Epidemiological and molecular studies on avian influenza viruses in Egypt. PhD Thesis. Faculty of Veterinary Medicine, Cairo University, Egypt.
- Kilany, W. H., A. Arafa, A. M. Erfan, M. S. Ahmed, A. A. Nawar, A. A. Selim, S. G. Khoulosy, M. K. Hassan, M. M. Aly, H. M. Hafez, and E. M. Abdelwhab (2010).** Isolation of highly pathogenic avian influenza H5N1 from table eggs after vaccinal break in

- commercial layer flock. *Avian diseases* 54:1115-1119.
- Kilany, W. H., E. M. Abdelwhab, A. S. Araf, A. Selim, M. Safwat, A. A. Nawar, A. M. Erfan, M. K. Hassan, M. M. Aly, and H. M. Hafez (2011).** Protective efficacy of H5 inactivated vaccines in meat turkey poult after challenge with Egyptian variant highly pathogenic avian influenza H5N1 virus. *Veterinary microbiology* 150:28-34.
- Li, Y., Z. Lin, J. Shi, Q. Qi, G. Deng, Z. Li, X. Wang, G. Tian, and H. Chen (2006).** Detection of Hong Kong 97-like H5N1 influenza viruses from eggs of Vietnamese waterfowl. *Arch. Virol.* 151:1615–1624.
- OIE (2009)** (World Organisation of Animal Health). Manual of diagnostic tests and vaccines for terrestrial animals, Part 2.3.4. [Internet]. [cited 2009]. Available from: [http://www.oie.int/eng/normes/mmanual/A\\_summary.htm](http://www.oie.int/eng/normes/mmanual/A_summary.htm)
- Palese, P., and M. L. Shaw (2007).** Orthomyxoviridae: The viruses and their replication, p. 1647-1689. In D. M. Knipe and P. M. Howley (ed.), in: *Fields Virology*, 5th ed. Lippincott Williams & Wilkins, Philadelphia.
- Peyre, M., H. Samaha, Y. J. Makonnen, A. Saad, A. Abd-Elnabi, S. Galal, T. Ettel, G. Dauphin, J. Lubroth, F. Roger, and J. Domenech (2009).** Avian influenza vaccination in Egypt: Limitations of the current strategy. *Journal of molecular and genetic medicine : an international journal of biomedical research* 3:198-204.
- Pillai, S. P. S., M. Pantin-Jackwood, H. M. Yassine, Y. M. Saif, and C. W. Lee (2010).** The high susceptibility of turkeys to influenza viruses of different origins implies their importance as potential intermediate hosts. *Avian Dis.* 54 (Suppl. 1):522–526.
- Promkuntod, N., C. Antarasena, and P. Prommuang (2006).** Isolation of avian influenza virus A subtype H5N1 from internal contents (albumin and allantoic fluid) of Japanese quail (*Coturnix coturnix japonica*) eggs and oviduct during a natural outbreak. *Ann. N. Y. Acad. Sci.* 1081:171–173.
- Safwat, M. A. (2007).** Studies on avian influenza virus in Egypt. Master Thesis. Cairo University, Cairo, Egypt.
- Smith, G. J. D., X. H. Fan, J. Wang, K. S. Li, K. Qin, J. X. Zhang, D. Vijaykrishna, C. L. Cheung, K. Huang, J. M. Rayner, J. S. M. Peiris, H. Chen, R. G. Webster, and Y. Guan (2006).** Emergence and predominance of an H5N1 influenza variant in China. *Proc. Natl. Acad. Sci. U.S.A.* 7:16936–16941.
- Spackman, E., D. A. Senne, T. J. Myers, L. L. Bulaga, L. P. Garber, M. L. Perdue, K. Lohman, L. T. Daum, and D. L. Suarez (2002).** Development of a realtime reverse transcriptase PCR assay for type A influenza virus and the avian H5 and H7 hemagglutinin subtypes. *J. Clin. Microbiol.* 40:3256–3260.
- Tian, G., S. Zhang, Y. Li, Z. Bu, P. Liu, J. Zhou, C. Li, J. Shia, K. Yu, and H. Chen (2005).** Protective efficacy in chickens, geese and ducks of an H5N1- inactivated vaccine developed by reverse genetics. *Virology* 341:153–162.
- Tonga S., Y. Lia, P. Rivailierb, C. Conrardya, D. A. Alvarez Castilloc, L. Chenb, S. Recuencod, J. A. Ellisonsd, C. T. Davisb, I. A. Yorkb, A. S. Turmelled, D. Moranc, S. Rogersa, M. Shia, Y. Taa, M. R. Weile, K. Tangf, L. A. Rowef, S. Sammons, X. Xub, M. Fracef, K. A. Lindbladeg, N. J. Cox, L. J. Andersona, C. E. Rupprechtd, and R. O. Donisb, (2012).** A distinct lineage of influenza A virus from bats; *proc Natl Acad Sci USA* 109(11):4269:4274.