

Avian Influenza in Egypt

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Aim

In this mini-review we have presented some ecological aspects of the isolated avian influenza viruses in Egypt since 2006, genetic characterization and the effect of some physical and chemical agents on their activity.

Abbreviations: HI: Hemagglutination Inhibition; HA: Hemagglutination; HPAI: Highly Pathogenic Avian Influenza Virus

Introduction

HPAI is a disease of global concern because of the threat posed to food security in regions that are dependent on poultry as a main source of protein and livelihood. An additional concern is that the H5N1 virus may mutate and cause a human influenza pandemic in which millions of human lives would be threatened [1-4].

Egypt experienced the disease since the first introduction of highly pathogenic Avian Influenza HPAI H5N1 in 2006. The virus widely extended in very short time and infected commercial production sectors and backyards [5,6].

Egypt is considered a hotspot for the evolution of a pandemic potential virus either via antigenic drift of the H5N1 to increase its adaptation to humans or H9N2 or through re assortment with other IAV subtypes, especially H3N2 virus.

Material and Methods

i. Genetic characterization of Egyptian H5N1 and H9N2 viruses-Analysis of the haemagglutinin (HA) and phylogenetic tree, identified.

ii. Chicken litter from a commercial broiler farm was provided.

iii. The effect of temperature and UV light on the infectivity of avian influenza virus (Samples were divided into 4 groups as follows: The first group was tested at 28 °C. The second group was tested at 37 °C. The third group was tested at 56 °C. The last group was subjected to UV light exposure at a density of 4-5 w/cm² (25 °C), which was the average density during the day at the study site.

iv. Treatment with disinfectants (The solution of disinfectant was added to half of the boxes of contaminated chicken litter and half of the controls).

v. Viability of the virus (The filtrate from each of the groups was inoculated into three 9-11 day-old chicken embryonated

eggs according to WHO protocol. After 48hrs of incubation, the allantoic fluid was harvested and hemagglutination (HA) and hemagglutination inhibition (HI) tests were used to identify the recovered virus).

Results and Discussion

Phylogenic and genetic changes of H5 n1 and H9n2 in Egypt

Conditions that favor genetic changes in the virus via antigenic shift (mutation) and drift (re assortment) are most likely to favor the emergence of HPAI viruses. The natural tendency for influenza viruses to regularly undergo point mutations, an inherent characteristic of RNA viruses, means that new viruses are continually being produced, with differences in genetic fitness [7]. Conditions that exert selective pressure on circulating viruses at both the host and population level act to increase the rate of mutation of viruses and therefore favour the appearance and establishment of dominant virus strains [8]. Thus, intensive poultry production systems in which a continuous and easily accessible source of susceptible hosts are present are considered prime conditions under which pathogenicity may emerge. Other cited conditions have been the inadequate use of vaccinations or incomplete vaccination coverage that has allowed field strains to reassort with vaccinal strains [9].

Genetic characterization of egyptian H5N1 viruses

Analysis of the haemagglutinin (HA) phylogenetic tree identified that the viruses are fall within the clade 2.2.1.1.

Genetic characterization of egyptian H9N2 viruses

Analysis of the haemagglutinin (HA) phylogenetic tree, identified that the viruses are fall within the A, B and C groups.

Meta-analysis situation and evaluation of the epidemic studies and their measurements of AIV in Egypt

It had wide variations according to Cochran's Q statistic and Higgins and Thompson's I², as shown in Figures 1 & 2 with low precision will show a wide variation in effects, while studies with high precision will show much less variation [10].

Meta-analysis situation and evaluation of the phylogenetic studies of AIV in Egypt

It had low variations according to Cochran's Q statistic and Higgins and Thompson's I², as shown Figures 1 & 2 with high precision will show much less variation [10].

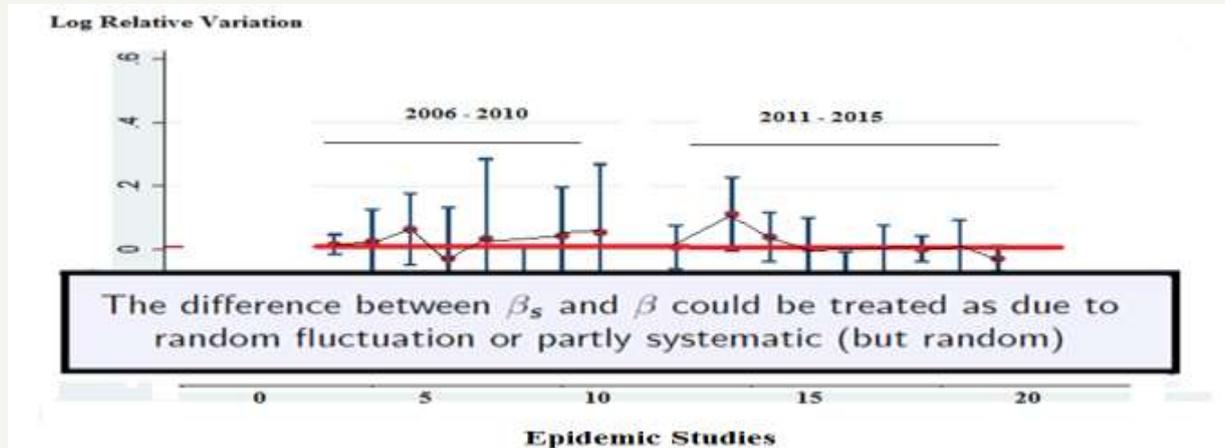


Fig.1: The studies had wide variations according to Cochran's Q statistic and Higgins and Thompson's I². The proportion of total variability explained by heterogeneity: Values > 25% so, it is High. (Phylogenetic analysis of the hemagglutinin gene of subtype H5N1 viruses: The H5N1 HPAI Egyptian viruses isolated in 2007 belong to clade 2.2.1, viruses isolated in 2008 to clade 2.2.1.1 while viruses isolated in 2010 belong to both sub-clades: 2.2.1 and 2.2.1.1, 2010 – 2014 sub-clade 2.2.1.1).

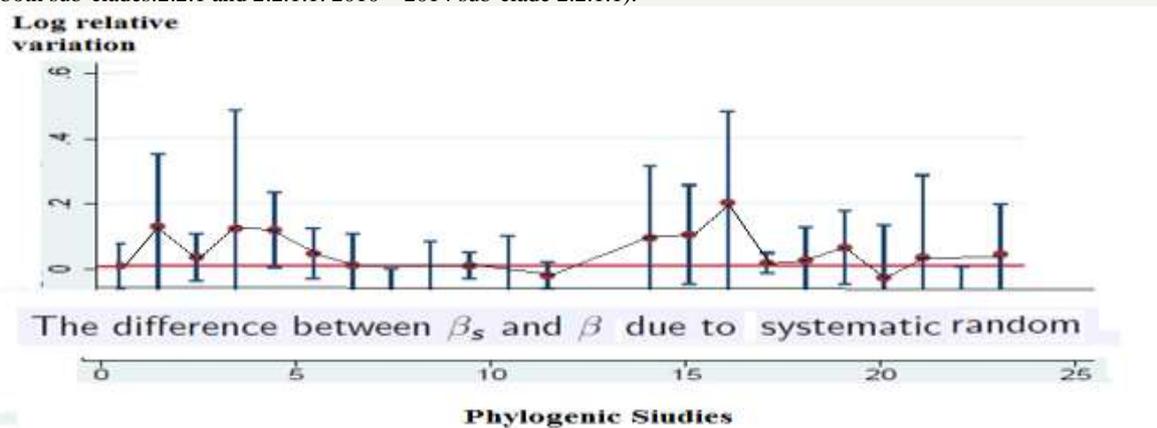


Fig.2: Meta-analysis situation and Evaluation of the phylogenetic studies of AIV in Egypt. The proportion of total variability explained by heterogeneity: Values < 25% so, it is low.

Kaoud et al. [11] have presented some aspects of the highly pathogenic avian influenza virus (HPAIV) in Egypt and the effect of some physical and chemical agents on its activity. The effect of temperature and UV light on the infectivity of isolated avian influenza virus H5N1 virus in litter could be inactivated by increasing temperature above 50 °C for at least 24hr. UV light could not destroy the infectivity of the virus completely even after exposure for 48 hr.

Disinfectants were evaluated in their study including Envirolyte, Virkon®-S 1%, Aldekol 0.5%, and bioscentry 0.5%. The results revealed that Envirolyte was very effective in reducing the titre of H5N1 virus after 10, 30min and 12hr of exposure at 25 °C from (28) to (23) but it completely destroying the virus after 24hr of exposure at 25 °C with complete reduction in HA activity. The

results also, revealed that Virkon- S® and Aldekol were effective in reducing the titre of virus after 30 min of exposure at 25 °C to (23) without any additional reduction afterwards for Virkon- S®. While, Aldekol succeeded in reducing the titre of H5N1 virus after 12hr of exposure at 25 °C to (22) but without any inactivation of HA.

Conclusion

The increase in human cases may be attributed to: increased circulation in backyard poultry, lower public health awareness of the risks, and seasonal factors such as closer proximity to birds due to cold weather. It is also suggested that co-circulation of A (H9N2) may be involved through multiple mechanisms in the enhanced spread of A(H5N1) amongst poultry in Egypt.

We think that avian influenza virus undergoes frequent antigenic drift, which is the accumulation of point mutations in the antigenic domain of the HA protein. As a result, viruses with a slightly changed antigenic structure emerge and can escape the host's acquired immunity, whether this immunity is acquired by natural infection or vaccination. Therefore, to maintain optimal protection by vaccination, the presently prevailing strains of influenza virus need to be included in each year's, influenza vaccine, requiring yearly reevaluation and frequent changes to the vaccine formulation.

For successful reduction of the negative impact on human health and associated economic and food security consequences, it will be essential to strengthen animal and human disease surveillance and disease control programmes. H5N1 virus can be inactivated in the poultry farms/ hatcheries using high temperature (e.g. 56 °C), using of disinfectants of the material to be disinfected. However, it may not be practically feasible for the farmers. Use of disinfectants seems more appropriate and practicable. Consequently there is no need to depopulate the poultry sheds after AIV outbreak for long period of time before arrival of new stock if disinfectants are used appropriately. The disinfection of surfaces also plays a role in controlling the spread of disease to other facilities. Mostly liquid disinfectants have been used for this purpose.

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