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RESEARCH PAPER

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Serological Investigation of infectious with avian influenza (H9N2) in broilers and human population in Abhar city

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Abstract

Influenza virus H9N2 Low Pathogenicity of the viruses, it was considered as widely spread low pathogen in poultry, however there are a few reports of human infection by this virus. This study was conducted to evaluate if such infection existence in workers of slaughter house and patients in Abhar hospital. This study, was conducted in Abhar poultry houses and Abhar hospital. Two hundred samples from poultry, 100 samples from patients with non-respiratory signs, 75 samples from patients with respiratory signs and 25 samples from poultry slaughterhouse workers was taken. To prevent false positive and negative results in HI test the sera were treated by trypsin-periodate and concentrated RBC respectively, and then serums was tittered by HI test.Our results indicated 80 percent of poultry flocks, 44 percent of poultry slaughterhouse workers was positive in HI test against H9N2 influenza. The high percentage of positive cases and higher titers in slaughterhouse workers indicate that the infectivity of the virus requires close contact with birds or their droppings.

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Introduction

Influenzavirusesare divided into four species of A, B, and C, based on M and NPproteins. Influenza virusAaffects birds, mammalssuch aspigs, horses, whales, andhumans. Based on thechange in HA and NAsurfaceglycoprotein, Influenza Avirusisclassified intothe16HAsubtypesand9NAsubtypes.

Allthesubtypes have been isolated from birds (Swayne, 2008).

Avianinfluenza viruses are classified into two groups, based pathogenicity, of on low and highpathogenicity. The Compatibility of these viruses withbirdsandrareincidence of the diseaseinbirds has introduced themas anatural reservoirof thevirus. These virusescanbreak down theinterspeciesbarrierand transmitted tomammalssuch horses, seals aspigs, andhumanscaused the disease incidence. Amongthese viruses, thehyperacuteinfluenza viruses of the H₅ and H7 and a low pathogenicH9virusareconsidered as new influenza pandemic candidatesdue to direct transmissionto humansandcause diseasein humans.

In 1999and then in2003, virusinfectionwithA/HK/1073/99(H9N2) andA/Hk/2108/03occurredin humans and since then the virus has been more considered as one of the candidates of new pandemic diseases.

Thestudies found that the internal genes of the virus are very similar to H5N1 virus isolated from humans in 1997. Also, The H9N2 virus that has taken the six internal genes from H5N1 virus has been found in a wide range of Asian birds.

In a study conducted by Cameron *et al* (2000) on H9N2 viruses is olated from birds of Germany, Iran, Pakistanand Saudi Arabia during the years 1998 to 1999 it was revealed that the virus had a close relationship with the humans' infective virus in Hong Kong (Cameron *et al.*, 2000).

The study conducted by Karimi*et al*(2005) shows the similarity of the H9N2 virus with the virus (H9N2)

A/Quail/Hk/G1/97(Karimi, 2005).

Ina study conducted byMoosaKhaniet al on12 cases of H9N2viruses in Iran, considering the similarity of the viruses to Pakistani viruses of A/Hk/1037/99 and A/quail/Hk/G1/97, it could be concluded thatthese virusescaninfecthumans,too. These allindicate the extent of the outbreak of H9N2 in Asian poultry and humans (Moosakhaniet al., 2010).

The study conducted byMatroskwitchet al (2000) showedthat the amino acidleucineat position of 226hemagglutininprotein are found specifically in humanstrains of H2 and H3, they also demonstrated that the conversion of glutamine toleucineat position of 226 of the virusA/Hk/1037/99 (H9N2) occurredinHong Kongwhichmakesidentifying the receptors(2, 6) of αacid silence in humans andhuman infection with this virus (Matrosovich et al., 2004).

In the study conducted by MoosaKhaniet al(2010) on 12 isolated H9N2virusesfrom Iran,it was found that100f12viruses hadleucine amino acid at position of 226 and the two other viruses hadglutamineamino acid(Moosakhani et al., 2010).

The mentioned 10 viruses were often related to 2005, indicating a changein the direction of the virus for human infection (Karimi, 2005). So, following theamino acidchangesat position of 226 of virus H9N2 andacquirethe ability toidentifytherecipienthuman flu, the virus canbeconsideredas acandidate for thedevelopment ofpandemic(Matrosovichet 2000). According to the conducted researches on the amino acids of Iranian H9N2virushemagglutininproteinbreakdown,the sequence of amino acids in this region is R-x-x-R (x is a non-basic amino acid, and R is arginine) that has thepotentialto becomeacutesequence.

Furthermore the studies conducted by Karimi, Torogi, and MoosaKhani*et al*confirmthesequences (Karimi, 2005- Moosakhani *et al.*, 2010). Withrespect tothe changes of 12 amino acids in

H9N2virusesisolated frompoultryinIranin 2010by thestudy of Moosa Khani, in case of the circulation of the virus in the Iran poultry farms, the incidence ofpointmutationsofthese virusesispossible(Moosakhani et al., 2010).

Materials and methods

A of200human bloodserum total samplesand200blood samplesfrom broilers prepared for slaughteringwere collected.

Humanbloodsamples of were consisted ofhospitalized 100serumsamples patients without respiratory symptoms (patients in the emergency department) was collected and 75 serum samplesofpatients admitted tohospital withrespiratory complicationsand25serum sampleswere collected fromslaughterhouse workers inAbhar.

The patientswithrespiratory symptomsandpatientswithoutrespiratory symptomswereidentifiedandseparatedandblood sampleswere takenfromthem.

Then, about 1ml of the considered patients' serum samples were obtained from the hospital laboratory in 1.5 ml tube and transmitted to microbiological lab of veterinary college of Tabriz in order to search flue antibody using HI method. The subtype antigen of commercial H9N2was used. 20blood samples from broilerflockwere each collected (a total of10broilerflocks) and aftertransferringthe samples to thelaboratory, the samples were centrifugedandtheserumwastested using HI method.

Serumdigestion using Trypsin - Heat - Periodate Treatment method

Toremovenonspecificinhibitorof hemagglutination (α, β, γ)inhuman serumandto avoidfalsepositivetest result, hemagglutination inhibition(HI)was performed. In short, 5.0volumesof trypsin were added 1volumeofserum(15.0mltrypsin to +3.0mlserum), then the trypsin containing serum was inactivated at 56°C water bathfor 30 min. Then it has beencooledto room temperature. 3volumesofo.011 M

of meta potassiumperiodate(0.9 ml) were addedto theserum. The mixture was incubated for 15 min at room temperature. After15minutes, 3volumes of 1%glycerol(0.9 ml) were added to the mixture. Then, the mixture was incubated again for 15minat room temperatureand finally 2.5 volumeof85% normal saline (0.75 ml) was added to the mixture until the serum was reached to a finaldilution of 1.10 (Who, 2005).

Detection of non-specific agglutinins in the digested serum

Toavoid the false negative result of hemagglutination inhibition, the nonspecificagglutinins should be detected in digestinghumanserum. In the case of a positive result, they must be absorbed. In short, aserial dilution of digested serums was madeina96hole U-shapedplate, then 1% chicken erythrocytes were added to the plate and they were incubated Aat temperature(22-25°C)for room 30minutes.Finally,ifcoagulum is formed afterthisperiod, the serumcanbe usedfortesting. However, in the case of agglutination, the nonspecific agglutinins should beabsorbedinserumbycondensed redblood cells in serum(Who, 2005).

Nonspecificagglutininabsorption thein digestedserumbycondensed red blood cells

Twentyvolumes of serum are addedto1 volumeof rinsed condensed red blood cellsand are mixed thoroughly. The mixture is placed at 4°C for an hour. Then the mixture is centrifugedat1200 RPM. The supernatant liquid wasremoved and tested for the presence of non-specific agglutinins. The practice will becontinued untilnegativeserum result is obtained(Who, 2005).

Statistical analysis

To study thedata of thestudythe Ttest, ANOVAandthe chi-square testwas usedbased onthe results of thestudy.

It should be noted that SPSS software (version 19) was used for statistical analysis of the data.

Results

The amounts of antibody in 10 poultry flocks were evaluated and the results of the farms with flu symptoms and non-flu farms (vaccinated) were evaluated using T-test and the

results are expressed in Table 1. The results of the study showed that there is a significant difference (p<0.01) between affected broiler farms and the farms without the flu.

Table1. Mean, standard error, standard deviation, and statistical significance of antibody titersagainstinfluenzain understudied poultryflocks.

Group	Mean±SE of antibody	SD	P value
Flocks with flu	5.97±0.179	2.26	_
Flocks without flu	0.85±0.192	1.11	0.001

Eight of10flocks(80%) had symptomsofthe avian influenzain thehemagglutination inhibitiontestsalsoshowedhightitersand2flocks(20%)

were unaffected and their antibody titer was due to vaccination whicharedescribed in Table 2.

Table2. Percentage of infected and uninfected bids in understudied flocks.

Group	Infected flocks percentage	Uninfected flocks percentage
Infected flocks	80	8
Uninfected flocks	20	2

Theantibody titerin samplesfromslaughterhouse workers, refferings to Abhar Hospitalemergency wardandhospitalpatients in the pulmonary wardwas evaluated. The results of the study showed that there is a statistically significant difference between the 3 study groups (p < 0.01). Also, the results of Duncan testshowed that the difference between antibody titer of slaughterhouse workers with the other two groups was significant (p < 0.05).

The results showed that 11 out of 25 samples (44%) obtained from the slaughter house workers had positive antibody titers against influenza and 4 out of 100 samples obtained from admitted reffering to the emergency ward (4%) were positive, and 8 out of 75 samples (10.66%) of the pulmonary ward patients had positive influenzatiters which are described in Table 3.

Table3. Mean, standard error, standard deviation, and statistical significance of antibody titersagainstavianinfluenzainthe understudied human population.

Group	Mean±SE	SD	P Value
Slaughterhouse workers	3.48±0.26*b	13	0.001
Emergency ward referring	1.31±0.11 ^a	1.13	
Pulmonary ward referring	1.31±0.21 ^a	1.27	

Table3. Percentage infected human population by poultry influenza.

Group	Infection percentage	Infected number/total samples
Slaughterhouse workers	44	11/25
Emergency ward patients	4	4/100
pulmonary ward patients	10.66	8/75

It should be noted that all positive casesofemergency andpulmonary wards have already worked ataviculture or had poultry farm.

Discussion and conclusion

Othersimilar studies in other parts of the world have been conducted toassess the prevalenceoftheH9N2virususingserologicalandvirolog icaltechniquesthat conform toobtain results in this study. In a study conducted by Chenget al (2002) todetermine the distribution of poultryinfluenzaA (H9N2)amongchicksandpeopleinShenzhenofChina, they isolated 27 tags of H9N2strainfrompoultry,but noviruswasisolatedfromhuman

cases. However, approximately 26% of the sera obtained frompatients and 7% of chickensera had H9N2 antibody, and HI antibodyinhuman serumhas beenfrom isolatedviruses from chickens (Cheng et al., 2002).

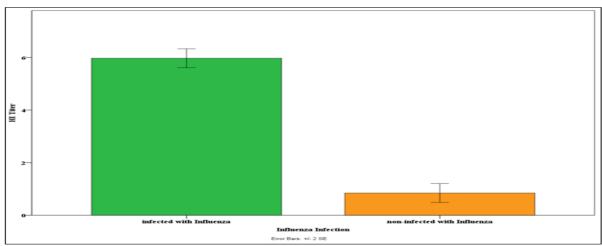


Fig. 1. Comparison of antibody titer meanagainst influenzain understudied flocks.

So, there is aclose relationship between H9N2 antibodies withvirusesisolated fromchickens.In another study, Li et al., in 2004investigated theoutbreak ofavian influenzaamong poultryandhumansinGuangzhou, China. Their study demonstrated a positiveH9N2antibodyin12.8% ofthe chickens's eraand 5.1 % of poultry farms and slaughterhouse workers. Also, they found that there was a close relationship betweenvirusesisolated fromchickenswithantibodiesagainst this virusin humans (Li et al., 2004).

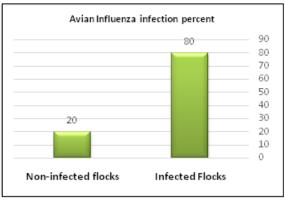


Fig. 2.Percentage of occurrenceor non-occurrence of influenzainunderstudied herds.

Inanother studybyGouet al.,in 2000,on a patient's serum inGuangzhou who was in recovery period and viruswasseparatedfromhis H9N2 serum. compare thelevels of antibodiesagainst thevirus in hismother andit was found thatthepatient had H9N2 titers at1.400-1.640.Also, his mother had antibodyagainstthe samevirusat 1.2 levels. It was likely thatthemother has been infected with the virus through contact withbirdsor throughairborne particles(Gouet al., 2000). Furthermore, Guoet al., in thestudyin 1999to determinewhether theH9N2viruscaninfect humans, took serum and throatsamplesfrompatients withflu symptoms, and chicken, and they found thatthe patientswerecarrying theH9N2virus.Also, they found thatabout 19% of them antibodiesagainst theH9N2virus have intheirserum(Guo et al., 1999).

It is revealed, by comparing theresults ofthesestudies withthepresent study, that the results are similar to each other. Interestingly, in these studies the isolated viruses from human viruses isolated from chickens that the poultry farm or slaughterhouse workers were in close contact with them were used as antigens for the HI test, and positive HI antibody of their serum has been

against these antigens. This indicates that in places such as slaughterhouses and poultry farms in which contact with the birds and their droppings is high, these viruses have a high capability to transmit and infect the humans and to compromise the body's immune system.

Theissue is similar to thehuman infectionwith H5N1 virus in which the most cases of humaninfection werein places where the poultry are infected with the virus and people indirect contact with poultry, became infected and sick.

Serological

studiesonclinicstaffandpoultryfarmsdemonstrated the positivity of HI antibody titers against the H9N2 virus (Momayez, 2000).

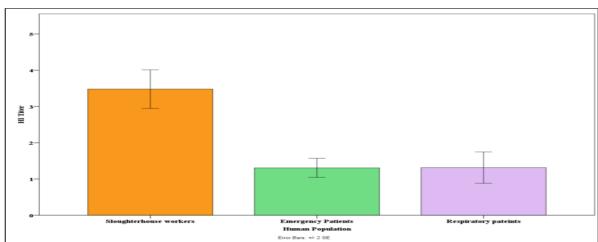


Fig. 3. Comparison of antibody titers against influenzain the understudied human population.

Inanotherstudy conducted on 200 ordinary people inshahre-Kord, it was found that about 16% of patients

hadpositiveserumHIantibodiesagainsttheH9N2 (Azizi and Aali, 2006). In these studies, the lack of understudied people's serum digestion by periodate or thereceiverdegradativeenzyme (RDE)to removenon-specificinhibitors ofhemagglutination, caused positive resultsinordinary peoplewhohadno contact withpoultry.It was likelydue toa false positivetestresult.

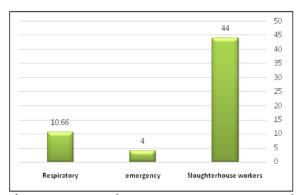


Fig. 3.Percentage of occurrenceor non-occurrence of influenza indifferent groups of humancases.

In the present study, the slaughterhouse workers and the hospital refferingssera were digested by trypsinperiodate-heat, which the resultsofbefore and afterdigestion were considerable; such that most of the pre-digested sera of the control group were positive in terms of H9N2 antibody. But, all serums pecimens were negative followed by digestion suggesting a false positive results and the presence of non-specific inhibitors of hemagglutination.

Surveys conducted byCameron(2000) andKarimi(2005) show the similarityofthevirusH9N2 of Iran with H9N2 (A/quail/Hk/G1/97 (virus infectinghumans inHongKong in1999) (Cameron *et al.*, 2000- Karimi, 2005).

The study byMayahiet al.,(2003) showed that the meanantibody titeragainstinfluenzaH9N2in people involved inpoultry industrywas significantly higher compared with(p <0.01)those not involved in the poultry industry. Also, theserumantibody titersof drivers and workers ofpoultrytransporttoslaughterhouses were significantly greater compared with the veterinarians(p<0.01) (Mayahi et al., 2003). The results of the study conducted by Goudarziet

al.,(2011) on thepeople workingin avicultures of the

East Azerbaijan provinceshowed that theantibody titer was negative in 790f96 cases(82%) and positive in

17 cases(18%) (Goudarzi et al., 2011).

Rahimianet al(2009) found that thehighestantibodytitersagainstH9N2 subtypewas inslaughterhouse workersthat19.7% of the testedserawere positive as well as in aviculture workers that 14.2% of the samples were positive. All the 29 serum samples ofFluVaccine preparation staffofRaziInstitutehad negativeantibody of subtypeH9N2.Only2%of those working inpoultryclinicshadpositivesamples (Rahimian et al., 2009).

The results of the present studyshowedthat44percentofworkers in the slaughterhouse had positive antibodies against influenz and the mean antibody titerwas 3.48 \pm 0.26. Four percentof

emergency patients and only 10.66% of pulmonary ward patients hadpositive antibodytiters againstinfluenza and all of which whom had a history of direct contact with poultry or working in a viculture. The results of this study are consistent with previous studies on the incidence avian influenza inhuman cases.

In a study conducted by Moosakhaniet al(2010) on12 cases of H9N2virusesisolated from Iran,it was found that10of12viruses had leucine amino acid at the position of226(similar tohumaninfluenza virusesH2N2andH3N2)and2otherviruseshad theamino acidglutamine(Moosakhaniet al., 2010).

Previous studies was indicating the changing of the virusesto infecthumans (Karimi, 2005). Therefore, theamino acidchangesat position226of virusH9N2andthe abilityto identify the human influenza receptor, the viruscan beconsideredas acandidate for thedevelopment ofpandemic(Matrosovichet al., 2000). So, thepositiveserologic testsinthis studyandother studies, suggestchanges in thereceptorbinding siteandthe abilityof the virustoinfecthuman.

Duetothe changes inthe amino acidsat positions226and 190 of thehemagglutininproteinofH9N2virus in Iran andspecificsequences of hemagglutininfracturezonesof these virusesand their capability to change into hyperacuteviruses. As well H9N2viruscirculatinginpoultry flocksandthose whoare in contactwithpoultryand occurrence some changestoinfecthumans, it is necessary tocombat thevirus.

In the present study the existence of antibodies specific for influenza A, H9N2 subtypein the serumofin fected people suggests the immune system response against the virus and indicating that infection occurs in them.

People working in the poultry industry maybe infected inhalation of airborneviruses or the consumption offoodcontaminated with viruses.

Sincethe symptoms ofinfluenza A,H9N2subtype in humansaresimilar to the symptoms of human influenza, such as, headache, fever, chills, muscle aches, cough, sore throat, sneezing, is therefore note a sily distinguishable fro meach other.

Conclusion

According to thementioned issues, slaughterhouses andavicultures workersshould bevaccinatedagainsthumaninfluenzaviruses in order to betteridentify and controlthe virusto prevent the confusion between human and avian influenza viruses.H9N2avianVaccineshould be made by new virusstrainsandit must be tried to make an effective and beneficial vaccine byrevisingthese strains, every 6 months. Also, all poultry flocks of the country must be vaccinated to prevent the circulation of the virus in the country's poultry.

Furthermore, the continuous study of the molecular prevalenceof the virusindifferent regions should be conducted in order to take care of thevirus changes topathogenesis andhumandisease. Theresults ofthe present study also indicate the infection of influenzaAsubtypeH9N2in people who are related to the poultry industry.

Considering the probability of influenza virus mutations, health organizations of the country need develop andimplement a comprehensive programtocontroland eradicatethevirusinthe poultry industry as well as people working inthe poultry industrymust be constantlymonitored for theirhealth.

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