ISOLATION AND CHARACTERIZATION OF PI-3 VIRUS FROM SHEEP AND GOATS

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225 nasal swab samples were collected from diseased sheep and goats for the isolation of PI-3 virus. The samples were taken from diseased animal at seven different Governorates Kaffer el sheikh, Alexandria, EL Behaira, Port Said, Demiatta El-Qalubia and Giza through the winter seasons of years 1999 to 2003. The virus was successfully isolated from three ovine samples after three successive passages on MDBK cells. The isolated viruses were then titerated and identified using different biological, serological and molecular assay.

Key words: Avian influenza H7, backyard, Egypt, RRT-PCR; ELISA, HI

.INTRODUCTION

Parainfluenza-3 (PI-3) is the most common virus infection of the respiratory tract of sheep and goats throughout the world. The virus belongs to family Paramyxoviridae of order Mononegaviralis, the non segmented negative single stranded RNA viruses (Pringle, 1991). The virus is generally

spherical and 150 to 350 nm in diameter but can be pleomorphic in shape. The genomic **RNA** contained within helical a nucleocapsid that is enclosed within the lipid bilayer. Inserted into the envelope are glycoprotein spikes that extended approximately 8 to 12 nm from the surface of the membrane (Kingsbury, 1991: Field, 1996).

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PI-3 virus was first isolated from sheep in 1966 (Hore, 1966) since then serological surveys have shown that this infection is widely spread in sheep population in many countries (Fenner et al., 1996). In Egypt, the first isolation of PI-3 virus from lambs was carried out in 1971(Baz 1971). The virus symptoms like mild revealed and nasal pyrexia, coughing discharge for several days and necropsy has shown area of pneumonia, particularly in ventral parts of the apical lobe (Hore and Stevenson, 1967)

The present study is directed to isolate the PI-3 virus from sheep and goats which naturally affected with respiratory manifestation, and to identify the isolated virus using several assays.

MATERIAL AND METHODS

Active surveillance

The surveillance was conducted in October, 2007 for three days, where 45 houses from 11 villages were included as shown in figure (1).

Samples

Serum samples: 207 serum samples were collected from

chickens (73), ducks (67), geese (52), pigeons (6) and turkey (9).

Tracheal and cloacal swabs: 200 tracheal swabs were collected from (66) chickens, (64) ducks, (47) geese, (12) pigeons, and (11) turkey and 200 cloacal swabs were collected from chickens (66), ducks (64), geese (47), pigeons (12), and turkey (11). As shown in table (1)

The birds were of different ages. The vaccination status against H5N1 virus was doubtful as told by the villagers, where two confirmed villages only were vaccinated at this time.

Avian Influenza Virus detection

Cloacal and tracheal swabs were collected from chickens, ducks, geese, pigeons and turkey. The swabs were immersed in 1-2ml viral transport media containing 2000000 I.U. penicillin, 2 mg streptomycin and 1000 I.U. mycostatin/ml. Briefly. five tracheal or Cloacal swabs were pooled together. RNA was extracted by using virus RNA Extraction Kit (QIAGEN, Valencia, Calif., USA).

The RNA were amplified using One-Step Real-time reverse transcription-PCR (QRT-PCR) kit (PG-Biotech; QIAGEN, Valencia,

Calif., USA) for detection of the matrix gene of influenza type A viruses.

Samples: 225 nasal swabs were collected from sheep and

goats suffering from respiratory manifestation at seven Govenorates through the winter seasons of years 1999 to 2003; the dilated data are shown in table (1).

Table (1) Number of nasal swab samples collected from sheep& goats in different localities in Egypt

Governorates	Localities	No. of sheep	No. of goats	Total number
Kaffer el sheekh	El karada	27	0	27
Alexandria		32	0	32
El Behaira	Wady el Ntron	2	0	2
Port Said		20	0	20
Demiatta		7	13	20
TO THE RESERVE	Meshthor	10	0	10
El- Qalubia	Tahanob	10	4	14
Giza	Abbatoir	55	0	55
	Oseem	16	4	20
	Nahia	11-11	14	25
Total nu	ımber	190	35	225

The samples were collected in sterile tubes containing 2 ml of sterile PBS and 200ul of stock antibiotic solution per each sample. The samples were taken on ice bucket. Clarification was done by centrifugation at 5000 rpm for 5 minutes at 4 c in cooling centrifuge. The supernatant was taken and kept at -70 c till used.

Viruses: first local bovine Egyptian PI-3 virus strain 45 was (kindly supplied from Rinder Pest like disease Department at Veterinary Serum and Vaccine production Institute Abbassia Cairo. Second bovine PI-3 virus strain SF₄ was kindly supplied from Rabies Department in VACSERA.

Cells: MDBK cells were kindly supplied from department of Virology, Fac. of Vet. Medicine Cairo University.

Heamagglutination test:

Samples were subjected to heamagglutination test to exclude the non-heamaglutinatinating samples from the isolation work. The test was conducted according to Singh& Baz (1966) using 1% G. pig RBC_s.

Isolation of PI-3 virus from HA positive samples:

The isolation was carried out according to Rulka (1987) using MDBK cells. Three successive passages were done for this isolation.

Titeration of isolates:

The harvest of the inoculated MDBK cells after three passages of the isolates was titrated. The test was performed according to Fray and Liess (1971) Briefly, in a microtiter plate 90ul of MEM free serum were dispensed into each well then 10ul of the suspected isolate suspension were added to the wells of the 1st column and serial ten fold dilutions were made.50.000 cells/ well were added while being in suspension. The plate was incubated at 37C in incubator with daily CO_2 microscopic examination. The TCID₅₀ was calculated according to Reed and Muench (1938).

Identification by heamadsorption technique:

The test was conducted according to Elizabeth et al., (1997). After 16 hours of inoculation of cells with isolates, the attached cells

were fixed for 30 minutes with formaldehyde in PBS (pH 7.2) then 1% suspension of G. pig RBC_s was added. The cells were examined under inverted microscope.

Identification of PI-3 virus by detecting inclusion bodies:

The test was made using Zinker Solution as a fixative according to Clayden (1971).

Identification of the isolates using Immunofluorescence:

The test was performed according to Vander Heide (1971) using cell culture staining chamber (CCSC). Each cup was inoculated with 100ul of the isolate and uninfected cups were included as control. The cups were overlaid with 100ul of anti PI-3 serum conjugated with FITC (supplied from Department of Virology, Faculty of Veterinary Medicin Cairo University) then examined under fluorescent microscope.

Identification of PI-3 suspected isolates by virus neutralization test:

The test was performed according to Murkami et al., (1983) Beta neutralization procedure using antibovine PI-3 hyperimmune serum (supplied from Serum& Vaccine Research Institute Abbassia, Cairo). The serum was

originally supplied from Imes, Iowa, USA).

Detection of PI-3 virus by polymerase chain reaction (PCR):

The genomic RNA was extracted from inoculated tissue culture by means of TRIZOL Reagent according to their manufacture instructions (commercially supplied by Life Technologies CO.) The TRIZOL Reagent is improvement to single step RNA

chromeznski and Sacchi (1987).

RT-PCR was employed using oligonucleotide primer pair designed from conserved site of fusion protein gene (F gene) as previously described by Lyon et al., (1997). The sequences of utilized primers as follow:

Sense (F)=5

CATTGAATTCATACTCAGCAC

3 Antisense (R) = 5 AGATTGTCGCATTT(AG)CCTC3

Table (2) Cycling protocol for amplification of F gene of PI-3

steps	Temp	Time	No. of cycle	
RT	47c	30 min.	One cycle	
Initial denaturation	94c	2 min.	One cycle	
Denaturation	94c	45 sec.	35 cycle	
Anealing	51c	45 sec.		
Extension	72c	1 min.		
Final extension	72c	5 min.	One cycle	
Preservation	4c	∞	and the second	

Analysis of the PCR product using agar gel electrophoresis.

The analysis was carried out according to Sambrook et al.,

(1989) using ethidium bromide stained agarose gel 1.5%

RESULTS

Results of heamagglutination test are presented in table (3). Five HA-

positive samples were cultivated and propagated on MDBK cells. Three passages in cell culture were applied and resulted three isolates which were designated as Oseem56/1999, Oseem184/2003 and Nahia70/2000 according to their localities and the year at which the samples were collected table (4) present the designation and passages results of the cell

culture work. The three isolates as well as local bovine strain 45 were titerated according to infectivity the results presents in table (5) It seem to be titers ranged from 10⁵ to 10^{6.6} TCID₅₀/ml with the highest test titer was given by the isolate Oseem56/1999 followed by isolate

Oseem 184/2003 and finally the isolate Nahia 70/2000. The isolated viruses were identified using hemadsorption, inclusion body detection, immunofluorescence, SNT as well as RT-PCR. Results of

Table (3) Haemagglutination titer of the five HA positive nasal swab samples and their localities.

Species .	Serial number	designated number	Locality	HA titer (HAU)
	1 0 30	56	Oseem	2 ⁶
	2	184		22
S	3	105	Port said	22
Sheep	4 .	101		23
	5	70	Nahia	24

Table (4) Results of cultivation of the HA positive (ovine) nasal swab samples

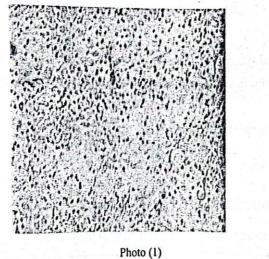
o. of passage	No. of positive samples (CPE)	No. of negative samples (no CPE)	Designated name of isolate
1 st passage	1	4	Oseem56/1999
2 nd passage	3	2	Oseem56/1999, Oseem184/2003 and Nahia70/2000
3 rd passage	3	2	Oseem56/1999, Oseem184/2003 and Nahia70/2000

CPE = cytopathic effect

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Table (5) Results of titration of PI-3 isolates on MDBK cells using infectivity test.

Designated name of isolate	Titer of virus (TCID50/ml)	
Oseem56/1999	10 ^{6.6}	
Oseem184/2003	10 ^{6.5}	
· Nahia70/2000	105	
PI-3 reference strain A ⁴⁵	10 ^{6.2}	



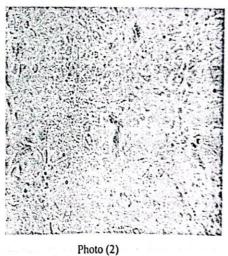


Photo (2) infected MDBK cells with the isolates after fixation and the positive heamadsorption.

Photo (3) normal non infected MDBK cells

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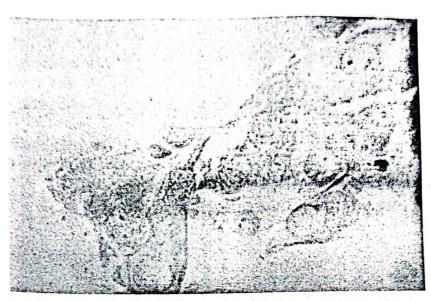


Photo (3) large esinophilic intracytoplasmic inclusion body

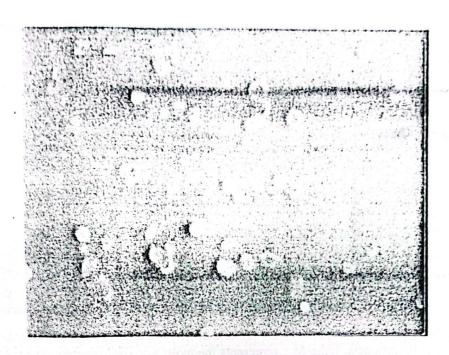


Photo (4) shows the observed intracytoplasmic perinuclear fluorescence in the in the positive FA-isolates on MDBK cells. Table (6) Results of virus neutralization test for PI-3 isolated viruses against antiparainfluenza-3 reference serum.

Photo (3) large esinophilic intracytoplasmic inclusion body.

Table (6) Results of virus neutralization test for PI-3 isolated viruses against antiparainfluenza-3 reference serum.

Isolate designated name	Neutralization titer on MDBK cell line
Oseem56/1999	512
Oseem184/2003	8
Nahia70/2000	256
(Reference A ⁴⁵ strain)	1024

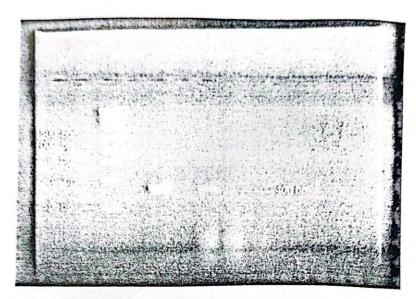


Photo (5) Shows the ethidium bromide stained 1.5% agarose gel electrophoresis of the PCR products. Lane M: represents the molecular DNA marker. Lane A: represents the amplified PCR product of approximately 400 bP, the conserved portion of F gene of PI-3 virus strain A⁴⁵ (positive control). Lane 1 to 3 represents the PCR products of the isolated viruses. Only Oseem56/1999 revealed the expected band (lane 1).

DISCUSSION

PI-3 virus causes, different clinical syndromes in sheep and specially goat respiratory infections (Ackermann and 2000). Serological Brodgen, surveys have shown that infection is widely spread among sheep population in many countries (Fishman, 1965; Pop et al., 1979:Brako et al. Saddour, 1987 and Riedmann et al., 1991). Many of serological data indicated that the ovine PI-3 virus is antigenically related to

bovine and human strains of PI-3 (Wood et al., 1975). In the presnt, work we tried to isolate PI-3 virus from naturally infected sheep and goats and RT-PCR was used as a recent tool for detection and identification of the isolated virus. The detection of PI-3 virus by heamagglutination, revealed five positive samples (all of them were from sheep samples representing 2.2% of the 225 collected samples. This low percentage of HA positive may be due to the high amount of virions (at least 105.5-10^{6.5} virions) required to give one work we tried to isolate PI-3 virus from naturally infected sheep and goats and RT-PCR was used as a recent tool for detection and identification of the isolated virus. The detection of PI-3 virus by heamagglutination, revealed five positive samples (all of them were from sheep samples representing 2.2% of the 225 collected samples. This low percentage of HA positive may be due to the high amount of virions (at least 10^{5.5}-10^{6.5} virions) required to give one HA unit (Redda 1964). The isolation was achieved through three passage applied on MDBK cells giving rise to three isolates developed a clear CPE. identification by detecting inclusion bodies revealed large esinophilic intracytoplasmic with no evidence to nuclear inclusions in contrast to bovine PI-3 virus that produced both cytoplasmic and nuclear inclusion similar results are previously reported (Howard et al., 1982). It was noticed that the three isolates were from lambs which may indicate be prevalence of PI-3 virus among sheep than in goats. However, the failure to isolate the virus from goat may be due to the lower number of collected samples from goats compared with those from

sheep and the fact that the animals were kept under relatively isolated conditions (Erasmus et al., 1967). The low titer of Oseem184/ 2003 isolate may be taken as criterion for heterogeneity (Gaul et al., 1982). This finding agreed with many of the previous studies, which have shown that at least three ovine PI-3 strains of different serotypes namely (G2 strain, CsL6 strain and DH-1 strain), do exist (Hore, 1966); St.George, 1969 and Howard et al., 1982). The heamadsorption used indicator system for the presence of PI-3 virus proved to be more rapid and sensitive than waiting until the cytopathic changes appear. This finding is similar to those described previously by St.George (1969) and Toth and (1990).RT-PCR Jankura technique waw performed in this work using two primers chosen from conserved site of fusion protein gene (F gene of whate PI-3 viruses). The primers were chosen in this site of genome as it is conserved among PI-3 viruses not specific for ovine PI-3 virus. Also, both sheep and goats can catch bovine PI-3 virus. Moreover, these primers could help us identification of the bovine PI-3 virus strain and the

standardization of the technique. Also, the same primers were developed by Lyon et al., (1997) and Maria et al., (1998). The detection and differentiation of PI-3 virus could be studied by amplification and sequencing of another part of the genome for

example HN gene Ruth et al., (1994), Echevarria et al., (2000) and Vecherov et al., (2003). Indeed, This study reports the isolation and characterization of parainfluenza-3 virus from sheep and goats in Egypt.

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