MOLECULAR CHARACTERIZATION OF EQUINE HERPES VIRUS -1 (EHV-1) IN EGYPT

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ABSTRACT

Equine Herpes Virus -1 is one of the most viruses affecting equide family (Horse, Donkey & Mule) causing severe economic losses to horse industry all over the world, due to respiratory manifestations, abortion of pregnant mares, and mayeloencephalopathy. Little reports have been made for investigation the prevalence and existence of equine herpes viruses (EHVs) in Egypt. In the present study, seventy clinical samples (6 placentas & 64 nasal swabs) collected from different governorate in Egypt. Virus isolation by inoculating tissue suspensions into MDBK cells revealed characteristic CPE for EHV after the fourth passage within 7 days from inoculation in the form of aggregation of the cells together, rounding, and finally cell detachment. Serological confirmation of the isolated sample by FAT revealed intracytoplasmic greenish yellow fluorescence. DNA extraction and PCR of suspected samples existed Successful amplification of 700 bp of UL45 gene of EHV-1 in 8 out of 70 examined samples.

Key words: Equine, Egypt, EHV-1, Virus isolation, FAT and PCR.
INTRODUCTION

Equine herpes viruses (EHVs) are significant causes lead to of sever illness and mortality in domestic horse population worldwide (Slater et al., 2006). Among the five recognized EHV, EHV-1 is enveloped with double stranded DNA genome, the genome was composed of a unique long (UL) region and a unique short (US) region, which was flanked by two inverted repeat regions (IR and TR) and encodes at least 72 unique genes and 4 diploid genes (Telford et al., 1992). Classified under family herpesviridea, subfamily alphaherpesvirinae, genus Varicellovirus (Slater, 2007).

EHV-1 was first isolated from aborted mare in Argentine in 1932 and 80% of horses had latent infection in the trigeminal ganglion and lymphocytes (Patel and Heldens., 2005). The involvement of lymphocytes results in development of a cell associated viraemia (Harless, 2006), which is responsible for the rapid dissemination of virus in secondary replication sites, including the gravid uterus spinal cord, testes and accessory sex glands without detectable effects on the fertility of stallions (Hebia et al., 2009).

Equine herpesviruses of type 1 and type 4 which are closely related antigenically and genetically, induce rhinopneumonitis and make long-live latent infections in the recovered horses in response to natural stresses as transportation, weaning and immune deficiency (Anonymous, 2008; Bresgen et al., 2012). The high prevalence is among young horses from weaning to 3 years of age (Walter, 2013).
EHV-1 is the main cause of abortions, paresis and neonatal foal deaths, due to necrotizing vasculitis and thrombosis resulting from infection of endothelial cells lining the blood capillaries. In these lesions, the initiation of infection is likely to be by reactivated EHV-1 from latently infected leukocytes. However, host factors responsible for reactivation remain poorly understood. Elderly horses, mares and febrile horses are at higher risk for developing severe neurological illness (Patel and Heldens., 2005).

EHV-1 induced neurological pathology is mainly restricted to the vascular epithelium of arteries supplying the central nervous system, although the trigeminal ganglion is also involved and the resulting inflammatory response leads to myeloencephalitis. (Goehring et al., 2006).

EHV-1 and EHV-4 are considered the most important pathogens both clinically and economically. Both viruses are primarily belonged to respiratory tract infections of varying degrees, and make long-live latent infections in the recovered horses. However, EHV-1 may progress to induce more serious symptoms in the form of abortion, still birth and neurological disorders (Allen and Bryans, 1986; Studdert et al., 2003).

Different methods have been used for EHV infections diagnosis in clinical samples (OIE, 2008). The virus isolation in cell culture derived from equine- has been the standard diagnostic approach. Apart from glycoproteins critical for virus egress to host cells, encodes several membrane proteins. Among them, UL45 is a class II protein dispensable for growth in vitro but was recently determined as an important virulence factor of HSV-1 neurotropism (Lucjan et al., 2008).
Further identification of the positive isolates was achieved by immunofluorescence, ELISA or immunoperoxidase using type-specific monoclonal antibodies (Borchers et al., 1997). During the last decades, several PCR assays were developed for identification and typing EHV (Dynon et al., 2001; Galosi et al., 2001; Varrasso et al., 2001; Szeredi et al., 2003; Diallo et al., 2006; Elia et al., 2006). The results of such assays well correlated with virus isolation in terms of accuracy and sensitivity, but surpassed it in simplicity, rapidity, time saving and the independence on the presence of infectious virus in the clinical sample (Varrasso et al., 2001).

MATERIAL AND METHODS

Sampling:

Table (1): Definition of samples used in the study.

<table>
<thead>
<tr>
<th>Disease manifestation</th>
<th>Sample collected</th>
<th>Animal age</th>
<th>Number of samples</th>
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| Respiratory symptoms  | Nasal swabs      | - Mature horses over 2 years  
|                       |                  | - Foals (6-18 months)        | 64 |
| Abortion              | Placenta         | Pregnant mares                  | 6  |

Seventy samples were randomly collected from Gharbia and Kafr - El Shekh Governorate, 20 out of them were apparently healthy and 50 were from the diseased horses and foals. The collected samples included swabs from different orifices in the body (nasal, pharyngeal and vaginal) in addition to tissue samples (placenta) from aborted mares (table 1). The specimens were grinded in phosphate buffered saline, pH 7.2, containing antibiotics of streptomycin, (50mg/ml), (2000 U/ml) penicillin, and (2 mg/ml) gentamycin. After freezing and thawing, the samples were clarified by centrifugation at 6000 rpm for 10 min; the collected supernatants were stored at -70 °C until used.
Virus isolation:

Isolation procedures were done according to OIE guidelines (OIE Terrestrial Manual 2013) Briefly, Use a T-75 flask of MDBK monolayer cells. The cells were grow under the standard culture conditions (37°C, 5% Co2 and 85% humidity) using an initial concentration 1.5x105 cells/ml. At 85% of cell confluence monolayer, tissue culture flasks are inoculated with 0.2 ml of the Prepared Samples. The monolayers are rinsed twice with phosphate buffered saline (PBS) After addition of 5 ml of supplemented maintenance medium (MEM containing 2% New born calf serum, the flasks are incubated at 37°C with daily inspection by microscopy for the appearance of characteristic EHV CPE then 3 cycles of Freezing and Thawing for Harvesting Supernatant is collected, stored in small aliquots at -80.

DNA extraction:

Viral DNA was extracted from different collected samples. Extraction was carried out according to the instruction of the DNA extraction kits QIA amp Viral DNA mini kits (Qiagen, USA). Briefly, about 130 μl from the samples was transferred to 570 μl of the Qiagen lysis buffer. Positive control extraction from Ab4p virus (Abortgenic strain of EHV-1) was kindly provided by Dr. A. J. Davison, Glasgow University- Scotland, the virus had a titer of 4x105PFU/ml, will run in parallel to each sample. DNAs were resuspended and stored at -70 ºC till use.

Polymerase Chain Reaction (PCR):

The DNA was further used for the amplification of full sequence of UL45 gene of EHV through gene specific forward 5'- (CTTGGA ATT CAT GGC AGG AGA CCC AAC A) -3' and reverse primer 5'-
(TTT GCG GCC GCT TAC CGG GGA AAC GGT ATG A) -3'. the reaction mixture consisted of 5 μl of extracted template DNA, 4.5 μl DD water, 1 μl of forward primer and 1 μl of reverse primer, 12.5 μl of PCR master mix. PCR thermo cycling using (T3 Biometra-Germany). The PCR programing was: initial denaturation at 94°C for 5min, followed by 5 cycles consists of 3 temp (98°C for 10 sec., annealing at 48°C for 30 sec and extension at 72°C for 30 sec., 30 cycles of 98°C for 10 sec., 58°C for 30sec., 72°C for 2 min, followed by1 cycle at 4°C for ∞. The PCR products were analyzed by electrophoresis on a 1.5% agarose gel containing ethidium bromide 0.5 μg/ml

**Immune Fluorescence Antibody Assay (IFA):**

Immune Fluorescence Antibody Assay was performed according to *Armin et al., (2004)*, briefly RK13 cells 40 -50% monolayer were infected with EHV-1 Ab4p virus at (MOI= 4) in 24 well plate at specific time point (overnight). The cells then fixed with 250 μl of ice cold 4% paraformaldehyde at RT for 15 min on bio shaker, after discarding of the paraformaldehyde, the cell were washed 3 times with 1X PBS. Adding 250 μl of 50mM NH4CL for each well and incubate at RT for 10 min on bio shaker, discard the NH4CL and add 240 μl of 0.2% triton 100 x for 10 min at RT on bio- shaker. Then the cells were blocked by 250 μl of 3% BSA/PBST for 30 min at RT. The cells running with 3 rounds of washing then incubated with 1st antibody (purified poly clonal antibodies prepared against UL45 gene which diluted 1/500 in 0.005% PBST for 1 h at 37°C in co2 incubator, washed three times by 0.005% PBST. The cells then incubated with the 2nd antibody FITC conjugated goat anti guinea pig IgG (sigma Aldrich, USA) diluted 1/200 in PBST for 1 h in 37°C incubator, after discarding the 2nd antibody, the cell were washed with 0.005 % PBST, and the slides were analyzed by fluorescence microscopy (Axiovert/Zeiss).
RESULT

Fig (1): Non inoculated MDBK monolayer cell culture as cellular control

Fig (2): MDBK cells inoculated with sample suspension showing cell rounding and detachment

Fig (3): Gel electrophoresis for PCR products of the examined samples. Lanes; 3-9 refer to the samples. Lane 1: Negative control, Lane 2: Positive control, M; 100 bp molecular weight DNA ladder.

Fig. (4): Fluorescence granules were observed after 24 hours post inoculated RK13 cells of the 3rd passage of inoculated samples using anti EHV-1UL45 conjugated with FITC
DISCUSSION

Worldwide, equine herpes viruses have a great economic and welfare impact on all horse sectors industry. They had several forms of the disease; including abortion, respiratory symptoms, and myeloencephalitis, that greatly affect the breeding and the performance ability of horses (Slater, 2007). In Egypt, there is a little information about epidemiology of EHV6s regarding the prevalence, circulating strains and factors predisposing the infection.

The presence of EHV was suspected from the epidemiological features of the collected samples. The study was continued for 2 years (2015 to 2016) using samples from all suspected cases either those showing sneezing, nasal discharge, coughing and/or abortion. Samples were inoculated in tissue culture cells (MDBK cells) which showed specific CPE in the form of rounding and cellular detachment after seven days of the 3rd cell passage. This may have been a consequence of the low virus titre in the inoculated samples. Later the virus became adapted to MDBK cells. Scott et al (1983) considered that EHV-1 exists in the infected mononuclear cells in non infective or subvirion forms. The obtained results were in agreement with Warda et al (2003) how succeeded in isolation of EHV-1 on BHK cells.

As a rapid monitoring of the EHV-positive samples, PCR was performed using specific primer sites against EHV-1UL45 gene, with an overall ratio of 11.4%. such assay achieve the advantages of specific and rapid identification of the viral antigen. Moreover, economically it provided a highly sensitive diagnosis tool of different EHV infection in a
large number of samples, the prevalence rate of the identified EHV-1 which determined in this report appeared to be lower than that recorded in other countries (Reubel et al., 1995; Nordengrahn et al., 2002). But in consistence with Haitham et al., (2011) who screening 93 clinical samples from Arabian horses for the presence of EHV antigens by semi-nested PCR assays and identified EHV-1, EHV-2, and EHV-4, with prevalence ratio 3.2%, 18.2%, and 7.5% respectively.

Immunological techniques (IF) using specific EHV-1 antisera showed fluorescence granules confirmed that the isolated virus was EHV-1. Isolation of the virus from nasal swabs and the time at which CPE was observed, suggest that the isolated virus is EHV-1 rather than EHV-4. Abdel Hafez et al., (2010) confirmed EHV-1 infection from fetal fluid obtained from aborted mares at 7th month of gestation using FAT.

The obvious role of EHV-1 in equine abortion is well documented throughout the world (van Maanen. 2002; Smith et al., 2003), although this virus strain was not recorded in any of the aborted samples which collected in this study.

REFERENCES


