### Research Article



### Pathological Identification of Egyptian Local Velogenic Newcastle Disease Virus Genotype VIId in Layer Chickens

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Abstract | Newcastle disease (ND) still threatening the poultry industry in Egypt despite stringent vaccination programs. This study was performed to evaluate the pathological lesions of Newcastle disease virus (NDV) genotype VIId in different organs of commercial layer chickens (ND antibodies-free). Therefore, forty, 1-day-old chicks were divided into 2 equal groups with 20 birds each. Group 1 served as the non-infected (negative control) group, while group 2 was infected at 28 days old by intranasal inoculation of 0.2 ml containing 106 EID<sub>50</sub> of NDV genotype VII. Tissue samples from trachea, lung, spleen and liver were collected from dead infected chickens at 4 days post-infection (dpi) and 6 dpi for detection of NDV RNA by real time Polymerase chain reaction (rRT-PCR). Tissue sections from lung, trachea, spleen, liver and proventriculus from the infected group and non-infected group were collected for histopathology at 5 dpi. The clinical signs in infected group appeared at 3 dpi and lasted for the 6th dpi represented by obvious greenish droppings at 5th dpi, leg paralysis and torticollis at the 4th and 5th dpi. Mortality was 10, 6 and 3 birds at 4 dpi, 5 dpi and 6 dpi respectively in infected group with total mortality rate 95%. Grossly, enlarged mottled spleen, severe hemorrhages on the tips of periventricular glands, severe muscle congestion, hemorrhagic ulcers on cecal tonsils and severe congestion with hemorrhages on the thymus gland at 4th and 5th dpi. Detection of NDV nucleic acid by rRT-PCR revealed highest detection rate (80% and 100%) in respiratory tissues (pooled trachea and lung), moderate detection rate (60% and 66.7%) in spleen and lowest detection rate (30% and 33.3%) in liver at 4 dpi and 6 dpi respectively. Histopathological investigation of different organs revealed pneumonia, tracheitis, spleenitis with lymphoid depletion, hepatitis and moderate proventriculitis. We conclude that our velogenic Newcastle disease virus (vNDV) strain is viscerotropic neurotropic pathotype exhibited high morbidity and mortality rates (95%) associated with immunosuppression in infected layer chickens. Detection of NDV RNA by rRT-PCR in our infected chicken organs emphasizing tissue tropism and distribution of vNDV in different organs, also suggesting the use of rRT-PCR in detection of NDV infected organs. further periodical investigations are needed to evaluate vNDV pathogenesis in chickens with considering the virus distribution pattern in different tissues.

Keywords | Experimental infection, Genotype VII, Histopathology, NDV, rRT-PCR

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#### **INTRODUCTION**

Newcastle disease virus (NDV) is one of the most harmful poultry pathogens threatening the poultry industry causing a highly contagious septicemic, fatal and destructive disease that attacks mainly chicken and turkeys (Khader et al., 2020). Since its first appearance in Indonesia

in1926 (Alexander, 2000) the NDV is still widespread in most countries worldwide with the only exception; The Oceania countries that showed a relative freedom from the disease (Dimitrov et al., 2017). In Egypt, NDV had been firstly identified in 1948 (Daubney and Mansy, 1948). The disease became widely disributed throughout the country and recorded as an endemic disease by the beginning of the

1960 (El-Nassary and Eskarous, 1960). According to their pathogenicity pattern, NDV is classified into velogenic (the most pathogenic), mesogenic (moderate pathogenic) and lentogenic strains (low pathogenic), while based on the tissue tropism and the developed clinical signs, the NDV is classified into viscerotropic, pneumotropic and neurotropic (Aldous and Alexander, 2001). High mortalities with nervous and intestinal pathology are recorded with viscerotropic velogenic and neurotropic phenotypes (Alexander et al., 2012; Nakamura et al., 2008). The main clinical symptoms of NDV are respiratory distress, diarrhea, and circulatory disturbance and in chronic cases, impairment of the central nervous system (Alexander et al., 1992). In chickens, the pathogenicity of NDV vary widely but mainly based on virulence of the virus. Other factors determine the pathogenicity of the disease such as the host species, age, immune status, infection with other organisms as well as dose, the route of administration and environmental stress. In some circumstances, infection with a virulent strain causes sudden deaths without major clinical signs in young chickens, while in adult birds the disease may be more protracted and less sever clinical signs such as respiratory distress including rales, sneezing, and wet eyes and/or nervous manifestations. (Cheville, 1972; Brown et al., 1999). NDV or avian paramyxovirus 1 (APMV-1) is a non-segmented, single-stranded, negativesense RNA virus in the family Paramyxoviridae (Phillips et al., 1998; de Leeuw and Peeters, 1999). Phylogenetically, based on the genome length and sequence of the F gene, NDV strains have been divided into classes I and class II (fifteen genotypes I-XV). Class I include avirulent strains and comprise a single genotype while class II contains 15 genetic groups including 10 previously established (I-IX, and XI) and five new genotypes (X, XII, XIII, XIV and XV). Class II also contains both virulent and avirulent vaccine strains such as LaSota and Hitchner B1 being used worldwide (Diel et al., 2012; Huang et al., 2004). Currently, the circulating strains from genotypes V, VI, and VII of class II are predominantly associated with disease outbreaks worldwide (Aldous et al., 2003). Genotype VII was initially divided into two sub genotypes: VIIa, representing viruses that emerged in the 1990 s in the Far East and spread to Europe and Asia, while VIIb, representing viruses that emerged in the Far East and spread to South Africa (Aldous et al., 2003). The two subgenotypes of VII are further divided into VIIc, d, and e, which represent isolates from China, Kazakhstan, and South Africa (Bogoyavlenskiy et al., 2009; Wang et al., 2006) in the same time, VIIf, g, and h, represent African isolates (Snoeck et al., 2009). In Egypt, the sub-genotype VIId is predominant causing several ND outbreaks in poultry (Radwan et al., 2013). NDV causes pathological changes in different organs which include tracheitis, pneumonia, pericarditis, myocarditis, and catarrhal

proventriculitis. Catarrhal enteritis, typhlitis, pancreatitis, perihepatitis, and nephritis have also been reported in NDV-infected poultry. Also Splenitis, atrophy of the Bursa of Fabricius, and encephalitis accompanied by lymphocyte and macrophage infiltration were also mentioned (Brar et al., 2017; Etriwati et al., 2017). The present study was designated to investigate the pathogenesis of NDV genotype VIId through recording the clinical signs, morbidity rate, mortality rate and pathological lesions with detection of histopathological alterations and virus distribution in different tissues according to r RT-PCR.

#### **MATERIALS AND METHODS**

#### SPECIFIC PATHOGEN FREE (SPF) EGGS

SPF Embryonated Chicken Eggs (ECE) obtained from Nile SPF (Koom Oshiem, Fayoum, Agriculture Research center- Ministry of Agriculture) was used for passage and titration of the isolated NDV isolate.

#### **EXPERIMENTAL CHICKS**

Forty 1 day- old male layer chicks obtained from commercial poultry company were used for pathogenicity assessment of NDV isolate (NDV/ chicken/FVCU/ Qalubia / Egypt/2019). Chicks were reared in previously cleaned and disinfected separate rooms on deep litter system with straw under optimal conditions from lightening, temperature and humidity. Birds were provided with commercial broiler plated (starter and grower) ration free from medication and provided with clean water ad-libitum. All procedures and bird handling were approved ethically by Institutional Animal Care and Use Committee (IACUC), Cairo University, Egypt (Approval number, CU/II/F/5/21).

#### NEWCASTLE DISEASE VIRUS (NDV) HEMAGGLUTINATION (HA) ANTIGEN

ND La Sota vaccinal strain were propagated in ECE and diluted to 4 HAU to be used as HA antigen in Hemagglutination inhibition (HI) testing of ND antibody.

#### NDV FOR PATHOGENICITY STUDY

Velogenic Newcastle disease virus (vNDV) local isolate (NDV/ chicken/FVCU/ Qalubia /Egypt/2019) isolated by authors during 2019 from 45-day old chicken flock suffered from respiratory manifestation and mortalities (5.4%) allocated in Qalubia governorate, molecularly characterized and submitted to genbank under accession numbers (MW172503). The used strain is related to NDV Genotype VIId.

#### INFECTIVITY TITRATION IN ECE

The propagated NDV suspension was titrated in [9-10] day old SPF ECE. 10-fold serial dilutions of the virus were prepared in saline solution contains antibiotic. Virus

suspensions were inoculated into 5 embryos for each dilution via allantoic sac (0.2 ml per egg). The inoculated embryos were incubated at 37°C and candled twice daily for 6 days. Dead embryos in first day were considered as nonspecific deaths. Slide haemagglutination (HA) test was applied on the allantoic fluid of inoculated chicken embryos to detect the HA-positive eggs. It was carried out according to the standard method described by (Anonymous, 1971; Grimes, 2002) and EID<sub>50</sub> was calculated according to Reed and Muench (Reed and Muench, 1938). The inoculation dose was 106 EID<sub>50</sub>/bird via intranasal instillation.

#### EXPERIMENTAL DESIGN

Forty 1 day- old male layer chicks were reared in disinfected separated rooms; feed and water were provided ad-libitum. Blood samples were randomly collected at one day- old, 14 day- old and 28 day of age (date of infection) from jugular vein of chickens (10 birds) (2 ml for each) on plain test tubes to separate serum to be used in HI test according to (OIE, 2012) using 1% freshly prepared chickens RBCs suspension in order to monitor the level of NDV maternal antibodies to detect the date of infection then chicks were divided into two groups (20 per each) as follow; control non infected group (group 1), infected group at 28 days old (group 2) were infected by intranasal route with 0.2ml of 106 EID<sub>50</sub> of NDV/ bird. Infected birds were observed daily for 10 days post infection (dpi) with recording of morbidity and mortality rates, clinical signs and pathological lesions (either macro or micro examination). Tissue samples from trachea, lung, liver and spleen were collected from dead infected bird to detect viral RNA in these tissues by rRT-PCR. Tissue sections from trachea, lung, liver, proventriculus and spleen were collected from all experimental groups at the 5th dpi for histopathological examination.

# NUCLEIC ACID (RNA) EXTRACTION AND RRT-PCR FOR DETECTION OF VIRUS NUCLEIC ACID IN DIFFERENT TISSUES

The viral RNA extraction was applied individually on the collected samples; spleen, liver and a pool of trachea and lung by Easy PureR Viral DNA/ RNA Kit (transbionovo, China) in accordance with manufacturer's instructions. Single step rRT-PCR assays using Trans ScriptR Probe One- Step qRT- PCR SuperMix (transbionovo, China) were conducted using specific oligonucleotide primers and probes for vNDV (Wise et al., 2004). The final reaction volume was 20  $\mu L$  including 4  $\mu L$  RNA template, 10  $\mu L$  2X TransStartR Probe qPCR mix, 0.4  $\mu L$  Trans ScriptR one- step RT/RI enzyme mix, 0.4  $\mu L$  of each of the forward and reverse primers, and 0.4  $\mu L$  probe together with 4.4  $\mu L$  nuclease free water. Thermocycling rRT-PCR conditions were 45°C for 5 min, 94°C for 30 sec, followed by 40 cycles at 94 °C for 5 sec and 30 sec at 54°C and

extension at 72°C for 10 sec.

#### HISTOPATHOLOGICAL EXAMINATION

Tissue sections from lung, trachea, spleen, liver and proventriculus were collected from the infected group and non-infected group and fixed on formalin 10%. The specimens were routinely processed in paraffin embedding method, sectioned, and stained with Hematoxylin and Eosin (H & E) for light microscopic examination according to Bancroft and Gamble (Bancroft and Gamble, 2008).

#### RESULTS AND DISCUSSION

#### MONITORING OF NDV SPECIFIC ANTIBODY TITERS

NDV mean antibody titers of sera collected at 1 day old, 14 days old and 28-day old chickens to detect the date of infection were  $6.00 \pm 0.63$ ,  $3.30 \pm 0.45$  and  $0.00 \pm 0.00$  respectively. At 28 days of age, birds were found to be free from any detectable antibody (sero negative chickens). (Values are expressed as mean of Log 2 $\pm$ SE).

#### CLINICAL SIGNS AND GROSS PATHOLOGY

Clinical signs begin at the 3<sup>rd</sup> dpi with marked depression with sleepy appearance, ruffled feather, greenish lose dropping and decrease in feed and water intake with no mortality. In 4th dpi: severe depression with sleepy appearance, foamy conjunctivitis, periorbital edema Figure 1A, neurological signs (leg paralysis) Figure 1B, marked decrease in feed and water intake and severe greenish lose dropping with sudden high mortalities Figure 1C. In the 5th dpi: obvious severe greenish lose dropping Figure 1D, neurological signs (torticollis) Figure 1E with severe loss of appetite and mortalities. 19 out of 20 birds in infected group died during 4th to 6th dpi Figure 2 while one bird in the infected group and all birds in the noninfected group remain healthy without obvious clinical signs and mortalities all over the observation period (10 dpi). Therefore, the morbidity rate was 19/20 (95%) also mortalities reach 95% in the infected group. The most prominent gross lesions of dead infected chickens observed during 4th and 5th dpi were enlarged mottled spleen Figure 3A, severe hemorrhages on the tips of periventricular glands Figure 3C, severe muscle congestion Figure 3B, hemorrhagic ulcers on cecal tonsils Figure 3D and severe congestion with hemorrhages on the thymus gland Figure 3E observed on postmortem examination.

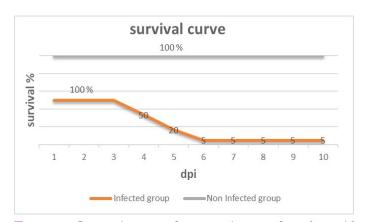
# DETECTION OF VIRAL NUCLEIC ACID (RNA) IN DIFFERENT TISSUES AT DIFFERENT INTERVALS POST INFECTION (DPI) USING RRT-PCR

Viral RNA detected by rRT-PCR in different tissues collected from dead infected birds at 4<sup>th</sup> and 6<sup>th</sup> dpi, the result revealed that (8/10) and (3/3) of the pooled trachea and lung tissue samples were positive at the 4<sup>th</sup> and 6<sup>th</sup> dpi,

respectively, (6/10) and (2/3) of spleen tissue samples were positive at the 4<sup>th</sup> and 6<sup>th</sup> dpi, respectively and (3/10) and (1/3) of liver tissue samples were positive at the 4<sup>th</sup> and 6<sup>th</sup> dpi, respectively Table 1.



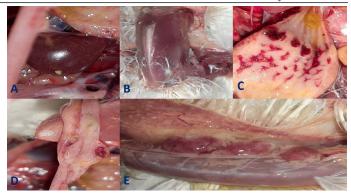
**Figure 1:** Clinical signs observed in the infected group at 4<sup>th</sup> and 5<sup>th</sup> days post- infection (dpi). A, foamy conjunctivitis and periorbital edema were observed at 4<sup>th</sup> dpi; B, neurological signs (leg paralysis) were observed at 4<sup>th</sup> dpi; C, sudden high mortalities were observed at 4<sup>th</sup> dpi; D, greenish lose droppings were observed at 5<sup>th</sup> dpi; E, neurological signs (torticollis) were observed at 5<sup>th</sup> dpi.



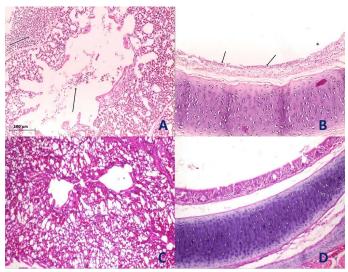
**Figure 2:** Survival curve after inoculation of 28 days old chickens with vNDV (infected group) and non-infected group of the experiment at different days post-infection (dpi).

**Table 1:** Detection of ND Viral RNA in different tissues at different days post infection (dpi).

dpi	Total number	Positive	
	of dead birds	Number	%
4 dpi	10	8	80
	10	3	30
	10	6	60
6 dpi	3	3	100
	3	1	33.3
	3	2	66.7
	4 dpi	of dead birds  4 dpi 10 10 10 6 dpi 3 3	of dead birds         Number           4 dpi         10         8           10         3           10         6           6 dpi         3         3           3         1



**Figure 3:** Gross pathological lesions observed in the infected group at 4<sup>th</sup> and 5<sup>th</sup> days post- infection (dpi) up on postmortem examination. A, Enlarged mottled spleen; B, severe muscle congestion; C, severe hemorrhages on the tips of periventricular glands; D, hemorrhagic ulcers on cecal tonsils; E, severe congestion with hemorrhages on the thymus gland.



**Figure 4:** Histological section of lung and trachea of the infected and non-infected group at 5<sup>th</sup> day post infection (dpi). A, lung showed moderate pneumonia with inflammatory exudate inside tertiary bronchi and secondary bronchi (arrow) with necrosed tertiary bronchi lining; B, trachea revealed tracheitis with noticeable necrosis of lining epithelium and mononuclear cells infiltration (arrow); C, normal lung (non-infected group) with no histopathological alterations; D, normal trachea (non-infected group) with no histopathological alterations.

#### HISTOPATHOLOGICAL EXAMINATION

Histopathological examination of the organs including lung, trachea, spleen, liver and proventriculus collected from the infected group and non-infected group at 5<sup>th</sup> dpi reveals that the lung showed moderate pneumonia with inflammatory exudate inside tertiary bronchi and secondary bronchi (arrow) with necrosed tertiary bronchi lining Figure 4A, trachea revealed tracheitis with noticeable necrosis of lining epithelium and mononuclear cells infiltration (arrow) Figure 4B, spleen showed spleenitis

with marked focal necrosis of lymphoid tissue (arrow) Figure 5A, liver showed hepatitis with mononuclear cells infiltration (arrow) and dilatation of hepatic sinusoids (yellow arrow) Figure 5B and Proventriculus suffered from moderate proventriculitis with mononuclear inflammatory cells infiltration within tunica muscularis (arrow) and epithelial cells degeneration Figure 5C. Non infected group showed no histopathological alterations in any examined tissue as illustrated in Figures 4C, 4D, 5D, 5E and 5F.

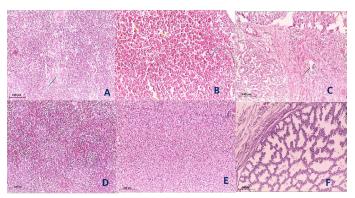


Figure 5: Histological section of spleen, liver and Proventriculus of the infected and non-infected group at 5<sup>th</sup> day post infection (dpi). A, spleen showed spleenitis with marked focal necrosis of lymphoid tissue (arrow); B, liver showed hepatitis with mononuclear cells infiltration (arrow) and dilatation of hepatic sinusoids (yellow arrow); C, Proventriculus suffered from moderate proventriculitis with mononuclear inflammatory cells infiltration within tunica muscularis (arrow) and epithelial cells degeneration; D, normal spleen (non-infected group) with no histopathological alterations; E, normal liver (non-infected group) with no histopathological alterations; F, normal Proventriculus (non-infected group) with no histopathological alterations.

During the last few years, severe NDV genotype VII outbreaks occurred in poultry flocks in Egypt causing severe economic losses in spite of stringent vaccination programs (Ewies et al., 2017).

Our experimental study is designated to evaluate the pathogenicity of a locally isolated vNDV strain (NDV/chicken/FVCU/Qalubia/Egypt/2019) (accession numbers MW172503) in commercial layer chickens. Chickens were infected via intranasal inoculation simulating the natural route of infection under field conditions (Alexander et al., 2012).

Our results revealed high morbidity and mortality rates (95%) in infected chickens indicating that our strain is velogenic, also molecular pathotyping of this strain through partial sequence of F gene including cleavage site motif showed that this strain is velogenic (unpublished data). Otherwise, two experimental studies were recorded using

different NDV strains of the same genotype (VIId) in boiler chickens, where one study recorded 72% mortalities at 25 days of age (Khader et al., 2020) and the other study recorded 80% mortalities at 28 days of age (Amer et al., 2018). All of these emphasizing the high morbidity and mortality rates associated with vNDV infection in chickens (Susta et al., 2011).

Notably, the clinical signs appeared at 3 dpi and lasted for the 6th dpi, which is likely attributed to the incubation period of ND (Miller and Koch, 2013). The clinical findings represented by obvious greenish droppings at 5th dpi, leg paralysis and torticollis at the 4th and 5th dpi indicate that our NDV isolate is a viscerotropic neurotropic strain. Similar findings were reported using different NDV strain of the same genotype in broilers (Khader et al., 2020; Mohammed et al., 2019; Mousa et al., 2019) and commercial male layer chickens vaccinated with Hitchner B1 and Lasota live ND vaccines (Amer, 2018). Also, the marked gross and histopathological lesions found in proventriculus confirm the viscerotropic tropism of our NDV strain. Similar findings were reported using different NDV strain of the same genotype in broilers (Mahmoud et al., 2019).

Although the used strains belong to the same genotype, the mortality rate in our infected layer chickens (95%) was higher than that recorded in boilers (Khader et al., 2020; Amer et al., 2018), also the onset of nervous signs was earlier in our infected layer chickens (4 dpi) compared to that recorded in boilers (Mohammed et al., 2019; Mousa et al., 2019), suggesting that layers possibly more vulnerable to velogenic neurotropic NDV infection (King, 1996). However, molecular fingerprinting is another factor to be investigated.

The necropsy of our infected chickens revealed severe systemic affections of muscles and internal organs like proventriculus, intestine and spleen emphasizing that our NDV strain is velogenic and able to replicate systemically in different organs (Eze et al., 2014). Similar findings were reported using different NDV strain of the same genotype in broilers (Mousa et al., 2019).

The marked gross affections in the gut associated lymphoid tissues including cecal tonsils and proventriculus gland as well as immune organs like spleen and thymus gland highlight the immunosuppressive effect of vNDV (Ezema et al., 2016). Similar findings were reported using different NDV strain of the same genotype in 30 day- old broiler chickens (Mousa et al., 2019).

detection of NDV RNA by rRT-PCR in spleen in our study suggesting that the gross and histopathological lesions of spleen are attributed to an NDV infection,

emphasizing the immunosuppressive effect of NDV (Ezema et al., 2016). Although the detection method was different, (Mousa et al., 2019) recorded similar detection rate (marked expression of viral antigen) in spleen of 30 day- old broiler chickens infected with different NDV strain of the same genotype by immunohistochemistry (IHC), emphasizing that rRT-PCR and IHC could be used for detection of NDV in infected organ (Hoffmann et al., 2009; Naglaa et al., 2014; Hussein et al., 2019).

Although mild respiratory signs, gross and histopathological lesions were recorded in our experiment, the highest detection rate of viral RNA was showed in respiratory organs (lung and trachea) at 4<sup>th</sup> and 6<sup>th</sup> dpi, this is basically because all NDVs initially replicate in respiratory system regardless this is because NDV replicates mainly at the portal of entry before it undergoes viraemia and disseminates into visceral organs (Wambura et al., 2006).

Although the detection method was different, (Khader et al., 2020) recorded similar detection rate in trachea and lung (highest viral density in trachea and lung at 5<sup>th</sup>, 6<sup>th</sup> and 7<sup>th</sup> dpi) of 25 day- old broiler chickens infected with different NDV strain of the same genotype by immunohistochemistry (IHC), also (Khader et al., 2020) recorded high viral RNA load in the trachea at 5<sup>th</sup> dpi by rRT-PCR. All of these emphasizing that rRT-PCR and IHC could be used for detection of NDV in infected organs (Hoffmann et al., 2009; Naglaa et al., 2014; Hussein et al., 2019).

Detection of NDV RNA by rRT-PCR at the 4<sup>th</sup> and 6<sup>th</sup> dpi and histopathological changes in liver of our infected chickens possibly attributed to an vNDV infection (Hussein et al., 2019). Similar histopathological findings were reported using different NDV strain of the same genotype in broilers (Aziz-ul-Rahman et al., 2019). Although the detection method was different, (Adi et al., 2012) reported similar detection rate (low viral density) in liver of chickens experimentally infected with viscerotropic vNDV using IHC method emphasizing that rRT-PCR and IHC could be used for detection of NDV in infected organ (Hoffmann et al., 2009; Naglaa et al., 2014; Hussein et al., 2019).

## CONCLUSIONS AND RECOMMENDATIONS

We conclude that our vNDV strain is viscerotropic neurotropic pathotype exhibited high morbidity and mortality rates (95%), gross and histopathological lesions in immune organs and gut associated lymphoid tissues causing immunosuppression in infected layer chickens. Also, detection of NDV nucleic acid by rRT-PCR in our

infected chicken organs was higher in respiratory tissues, moderate in spleen and lower in liver emphasizing tissue tropism and distribution of vNDV, also suggesting the use of rRT-PCR in detection of NDV infected organs. further periodical investigations are needed to evaluate vNDV pathogenesis in chickens with considering the virus distribution pattern in different tissues.

#### **NOVELTY STATEMENT**

This manuscript provides a data about clinico pathological picture and virus distribution in different tissues of infected layer chicken using new isolate of NDV related to genotype VII that is contributed in occurrence of severe outbreaks in poultry farms in Egypt.

#### **AUTHOR'S CONTRIBUTION**

All authors contributed in practical part (NDV isolation and infectivity titration, evaluation of experimental infection in layer chicken with NDV isolate. Also, all authors contributed in scientific writing and revision of this manuscript.

#### CONFLICT OF INTEREST

The authors have declared no conflict of interest.

#### REFERENCES

- Adi AAAM, Kardena IM, Astawa NM, Matsumoto Y (2012). Immunohistochemical detection of viral antigen in tissue of chickens experimentally infected with Newcastle disease virus. J. Vet., 13(3): 278-283.
- Aldous EW, Alexander DJ (2001). Detection and differentiation of Newcastle disease virus (avian paramyxovirus type 1). Avian Pathol., 30(2): 117-128. https://doi. org/10.1080/03079450120044515
- •Aldous EW, Mynn JK, Banks J, Alexander DJ (2003). A molecular epidemiological study of avian paramyxovirus type 1 (Newcastle disease virus) isolates by phylogenetic analysis of a partial nucleotide sequence of the fusion protein gene. Avian Pathol., 32(3): 237-255. https://doi.org/10.1080/0307945031000097831
- Alexander DJ (2000). Newcastle disease and other avian paramyxoviruses. Rev. Scient. Techn. Office Int. Epizooties, 19(2): 443-455. https://doi.org/10.20506/ rst.19.2.1231
- Alexander DJ, Aldous EW, Fuller CM (2012). The long view: a selective review of 40 years of Newcastle disease research. Avian Pathol., 41(4): 329-335. https://doi.org/10 .1080/03079457.2012.697991
- Alexander DJ, Campbell G, Manvell RJ, Collins MS, Parsons G, McNulty MS (1992). Characterisation of an antigenically unusual virus responsible for two outbreaks of Newcastle disease in the Republic of Ireland in 1990. Vet. Record, 130(4): 65-68. https://doi.org/10.1136/vr.130.4.65
- Amer MM (2018). Studies on pathogenicity of local Newcastle disease genotype VIIj and avian influenza H9N2 isolates

- to commercial vaccinated male layer chickens. Egypt. J. Vet. Sci., 49(2): 119-133. https://doi.org/10.21608/ejvs.2018.4958.1042
- Amer MM, Ahmed HM, Elbayoumi KM (2018). Pathogenicity
  of local identified NDV strain related to genotype VII to
  28-day old commercial broiler chickens. MOJ Bioequiv.
  Availab., 5(4): 227-230. https://doi.org/10.15406/
  mojbb.2018.05.00107
- Anonymous (1971). Methods of examining poultry biologics and identifying and quantifying avian pathogens. National Acad Sci., Washington DC.
- Aziz-ul-Rahman, Rohaim MA, El-Naggar RF, Mustafa G, Chaudhry U, Shabbir MZ (2019). Comparative clinicopathological assessment of velogenic (sub-genotype VIIi) and mesogenic (sub-genotype VIm) Avian avulavirus 1 in chickens and pigeons. Avian Pathol., 48(6): 610-621. https://doi.org/10.1080/03079457.2019.1648751
- Bancroft JD, Gamble M (2008). Theory and practice of histological techniques. Elsevier Health Sciences.
- •Bogoyavlenskiy A, Berezin V, Prilipov A, Usachev E, Lyapina O, Korotetskiy I, Zaitceva I, Asanova S, Kydyrmanov A, Daulbaeva K, Shakhvorostova L (2009). Newcastle disease outbreaks in Kazakhstan and Kyrgyzstan during 1998, 2000, 2001, 2003, 2004, and 2005 were caused by viruses of the genotypes VIIb and VIId. Virus Genes, 39(1): 94-101. https://doi.org/10.1007/s11262-009-0370-1
- •Brar RS, Leishangthem GD, Gadhave PD, Singh ND, Banga HS, Mahajan V, Sodhi S (2017). Diagnosis of the Newcastle disease in broiler by histopathology and immunohistochemistry. Indian J. Vet. Pathol., 41(1): 60-62. https://doi.org/10.5958/0973-970X.2017.00014.1
- Brown C, King DJ, Seal BS (1999). Pathogenesis of Newcastle disease in chickens experimentally infected with viruses of different virulence. Vet. Pathol., 36(2): 125-132. https://doi. org/10.1354/vp.36-2-125
- Cheville NF (1972). Pathogenesis of virulent Newcastle disese in chickens. J. Am. Vet. Med. Assoc., 161: 169-179.
- Daubney R, Mansy W (1948). The occurrence of Newcastle disease in Egypt. J. Comp. Pathol. Therapeut., 58: 189-200. https://doi.org/10.1016/S0368-1742(48)80019-6
- de Leeuw O, Peeters B (1999). Complete nucleotide sequence of Newcastle disease virus: evidence for the existence of a new genus within the subfamily Paramyxovirinae. J. Gen. Virol., 80(1): 131-136. https://doi.org/10.1099/0022-1317-80-1-131
- Diel DG, da Silva LH, Liu H, Wang Z, Miller PJ, Afonso CL (2012). Genetic diversity of avian paramyxovirus type 1: Proposal for a unified nomenclature and classification system of Newcastle disease virus genotypes. Infect. Genet. Evolut., 12(8): 1770-1779. https://doi.org/10.1016/j.meegid.2012.07.012
- Dimitrov KM, Afonso CL, Yu Q, Miller PJ (2017). Newcastle disease vaccines. A solved problem or a continuous challenge? Vet. Microbiol., 206: 126-136. https://doi. org/10.1016/j.vetmic.2016.12.019
- El-Nassary BB, Eskarous JK (1960). The distribution of fowl plague and Newcastle disease in upper Egypt. Arch. Mikrobiol., 36(2): 147-150. https://doi.org/10.1007/ BF00412282
- Etriwati DR, Handharyani E, Setiyaningsih S (2017).
   Pathology and immunohistochemistry study of Newcastle disease field case in chicken in Indonesia. Vet, World, 10(9): 1066. https://doi.org/10.14202/vetworld.2017.1066-1071

- Ewies SS, Ali A, Tamam SM, Madbouly HM (2017). Molecular characterization of Newcastle disease virus (genotype VII) from broiler chickens in Egypt. Beni-Suef Univ. J. Basic Appl. Sci., 6(3): 232-237. https://doi.org/10.1016/j. bjbas.2017.04.004
- Eze CP, Okoye JOA, Ogbonna IO, Ezema WS, Eze DC, Okwor EC, Ibu JO, Salihu EA (2014). Comparative study of the pathology and pathogenesis of a local velogenic Newcastle disease virus infection in ducks and chickens. Int. J. Poult. Sci., 13(1): 52. https://doi.org/10.3923/ijps.2014.52.61
- Ezema WS, Eze DC, Shoyinka SVO, Okoye JOA (2016).
   Atrophy of the lymphoid organs and suppression of antibody response caused by velogenic Newcastle disease virus infection in chickens. Trop. Anim. Hlth. Prod., 48(8): 1703-1709. https://doi.org/10.1007/s11250-016-1147-x
- Grimes SE (2002). A basic laboratory manual for the smallscale production and testing of I-2 Newcastle disease vaccine. RAP Publication, pp. 136.
- •Hoffmann B, Beer M, Reid SM, Mertens P, Oura CA, van Rijn PA, Slomka MJ, Banks J, Brown IH, Alexander DJ, King DP (2009). A review of RT-PCR technologies used in veterinary virology and disease control: sensitive and specific diagnosis of five livestock diseases notifiable to the world organisation for animal health. Vet. Microbiol., 139(1-2): 1-23. https://doi.org/10.1016/j.vetmic.2009.04.034
- Huang Y, Wan HQ, Liu HQ, Wu YT, Liu XF (2004). Genomic sequence of an isolate of Newcastle disease virus isolated from an outbreak in geese: A novel six nucleotide insertion in the non-coding region of the nucleoprotein gene. Arch. Virol., 149(7): 1445-1457. https://doi.org/10.1007/s00705-004-0297-8
- •Hussein EA, Hair-Bejo M, Omar AR, Arshad SS, Hani H, Balakrishnan KN, Yakubu Y, Saeed MI, Aini I (2019). Velogenic newcastle disease virus tissue tropism and pathogenesis of infection in chickens by application of in situ PCR, immunoperoxase staining and HE staining. Microbial. Pathog., 129: 213-223. https://doi.org/10.1016/j.micpath.2019.02.017
- Khader MA, El-Kady MF, Shaheed IB (2020). Viral distribution of Newcastle disease virus genotype VII in different organs of broiler chickens. Braz. J. Vet. Pathol., 13(2): 510-518. https://doi.org/10.24070/bjvp.1983-0246.v13i2p510-518
- King DJ (1996). Influence of chicken breed on pathogenicity evaluation of velogenic neurotropic Newcastle disease virus isolates from cormorants and turkeys. Avian Dis., pp. 210-217. https://doi.org/10.2307/1592391
- Mahmoud NK, El-Deeb AH, Emara MM, Abd El-Khaleck MA, Hussein HA (2019). Genotypes II and VIIdbased inactivated Newcastle disease vaccine reduces virus shedding. Virus Dis., 30(3): 453-461. https://doi. org/10.1007/s13337-019-00537-2
- •Miller PJ, Koch G (2013). Newcastle disease. Dis. Poult., 13: 89-138.
- •Mohammed FF, Mousa MR, Khalefa HS, El-Deeb AH, Ahmed KA (2019). New insights on neuropathological lesions progression with special emphasis on residence of velogenic Newcastle disease viral antigen in the nervous system of experimentally infected broiler chickens. Explor. Anim. Med. Res., 9(2): 145-157.
- Mousa MR, Mohammed FF, Khalefah HS, El-Deeb AH, Ahmed KA (2019). Comparative serological, histopathological and immunohistochemical evaluation of immune status of broiler chickens experimentally infected



- with velogenic Newcastle disease virus in different ages. Int. J. Vet. Sci., 8(3): 143-150.
- Naglaa M, Hagag ZA, Soliman MA, Saied M, Arafa HM (2014). Phylogenetic and histopathological characterization of Newcastle disease viruses reveals high prevalence of very virulent strains of genotype VIId in Egypt. Assist. Vet. Med. J., 61: 168-178.
- •Nakamura K, Ohtsu N, Nakamura T, Yamamoto Y, Yamada M, Mase M, Imai K (2008). Pathologic and immunohistochemical studies of Newcastle disease (ND) in broiler chickens vaccinated with ND: Severe nonpurulent encephalitis and necrotizing pancreatitis. Vet. Pathol., 45(6): 928-933. https://doi.org/10.1354/vp.45-6-928
- OIE (2012). Newcastle disease. Manual of diagnostic tests and vaccines for terrestrial animals. Pp. 555–574.
- Phillips RJ, Samson ACR, Emmerson PT (1998). Nucleotide sequence of the 5'-terminus of Newcastle disease virus and assembly of the complete genomic sequence: Agreement with the "rule of six". Arch. Virol., 143(10): 1993-2002. https://doi.org/10.1007/s007050050435
- Radwan MM, Darwish SF, El-Sabagh IM, El-Sanousi AA, Shalaby MA (2013). Isolation and molecular characterization of Newcastle disease virus genotypes II and VIId in Egypt between 2011 and 2012. Virus Genes, 47(2): 311-316. https://doi.org/10.1007/s11262-013-0950-y
- Reed LJ, Muench H (1938). A simple method of estimating fifty per cent endpoints. Am. J. Epidemiol., 27(3): 493-497. https://doi.org/10.1093/oxfordjournals.aje.a118408

- •Snoeck CJ, Ducatez MF, Owoade AA, Faleke OO, Alkali BR, Tahita MC, Tarnagda Z, Ouedraogo JB, Maikano I, Mbah PO, Kremer JR (2009). Newcastle disease virus in West Africa: new virulent strains identified in non-commercial farms. Arch. Virol., 154(1): 47-54. https://doi.org/10.1007/s00705-008-0269-5
- Susta L, Miller PJ, Afonso CL, Brown CC (2011). Clinicopathological characterization in poultry of three strains of Newcastle disease virus isolated from recent outbreaks. Vet. Pathol., 48(2): 349-360. https://doi. org/10.1177/0300985810375806
- •Wambura P, Meers J, Spradbrow P (2006). Determination of organ tropism of Newcastle disease virus (Strain I-2) by virus isolation and reverse transcription–polymerase chain reaction. Vet. Res. Commun., 30(6): 697-706. https://doi.org/10.1007/s11259-006-3291-7
- Wang Z, Liu H, Xu J, Bao J, Zheng D, Sun C, Wei R, Song C, Chen J (2006). Genotyping of Newcastle disease viruses isolated from 2002 to 2004 in China. Ann. New York Acad. Sci., 1081(1): 228-239. https://doi.org/10.1196/annals.1373.027
- Wise MG, Suarez DL, Seal BS, Pedersen JC, Senne DA, King DJ, Kapczynski DR, Spackman E (2004). Development of a real-time reverse-transcription PCR for detection of Newcastle disease virus RNA in clinical samples. J. Clin. Microbiol., 42(1): 329-338. https://doi.org/10.1128/JCM.42.1.329-338.2004

