ORIGINAL RESEARCH

Isolation and Molecular Detection of Marek's Disease Virus from Outbreak Cases in Chicken in South Western Ethiopia

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Background: Marek's disease virus is a devastating infection, causing high morbidity and mortality in chickens in Ethiopia.

Methods: The current study was conducted from March to November, 2021 with the general objective of performing antemortem and postmortem, isolation, and molecular detection of Marek's disease virus from outbreak cases in southwestern Ethiopia. Accordingly, based on outbreak information reported from the study sites namely, Bedelle, Yayo, and Bonga towns in southwestern Ethiopia, 50 sick chickens were sampled. The backyard and intensive farming systems of chickens were included in the sampling and priorities were given for chickens that showed clinical signs that are characteristics of Marek's disease.

Results: By clinical examinations, paralysis of legs and wings, gray eye, loss of weight, difficulty in breathing, and depression were recorded on all chickens sampled for this study and death of diseased chickens was observed. In addition, enlargement of the spleen and gross lesions of the liver and heart were recorded during postmortem examination. The death of infected chickens was observed in both vaccinated and non-vaccinated flocks. Out of 50 pooled feather follicle samples, Marek's disease virus was isolated from 14/50 (28%) by cell culture method and out of six tissue samples, the virus was isolated from 5/6(83.30%). By Real time polymerization chain reaction technique, which was targeted to detect the Meq gene, Marek's disease virus was detected from 18/50 feather follicles which accounts for 36% of sampled chickens.

Conclusion: In general, current study showed that the circulating Marek's disease virus in southwestern Ethiopia was caused by the oncogenic Gallid herpesvirus-2 (Serotype-1). Further research on molecular characterization of revolving virus in current and other regions is recommended for effective control of the disease through vaccination.

Keywords: Ethiopia, Marek's diseases, isolation, molecular detection, outbreak

Introduction

Chicken production is an essential agricultural activity in Ethiopia. The majority of the farmers and many others, keep at least a few chickens, which can play a significant role in nutrition, poverty alleviation, and food security. The country shared about 60% of the total chickens in East Africa. This number consists of all breeds of exotic, hybrid, and local. In detail, the estimation of the chicken population was about 56.53 million and with regard to breed, 94.3%, 3.21%, and 2.49% of the total poultry population are indigenous, hybrid, and exotic, respectively.³

The majority of chicken types reared in Ethiopia are local ecotypes, which demonstrate a large variation in body position, plumage color, comb type, and productivity¹⁰ and are managed under an extensive management system.²⁵ Village chickens have been reared as a source of family protein and income for many years and they have contributed to the country's economy. However, the overall chicken production system in Ethiopia is commonly known as it is not for the purpose of commercial targeted, decreased input, scavenging, and outdated management system entails domestic breeds. Accordingly, the economic importance of the sector is not still proportional to the large chicken numbers.¹⁷

In addition, the Newcastle disease, Infectious Bursal disease, and Marek's disease are making worse constraints on Ethiopia's poultry production. Among the major constraints of chicken productions, Marek's disease (MD) is a devastating infection, which replicates in lymphocytes, made the latent infection in CD4 (+) T lymphocytes,¹² and occurred in chickens by herpesvirus, which is scientifically known as *Mardivirus* genus.

This genus is classified into three serotypes namely; Gallid herpesvirus 2 (serotype 1), Gallid herpesvirus 3 (serotype 2), and Meleagrid herpesvirus 1 or herpesvirus of turkeys (HVT) (serotype 3). Out of the MD serotypes, the first one is virulent and a small number of them were the attenuated vaccine strains. The second species/serotype is naturally recognized as avirulent and the HVT also served as a vaccine, further identified as being a recombinant viral vaccine vector recently.²⁴

The disease, introduced in Ethiopia with the exotic breeds from outside, has now become a fear for backyard and commercial farming systems.¹⁶ Epidemiologically, MD is extensively distributed and causing the high morbidity and mortality of chickens at small-scale farming and backyard poultry farms in Ethiopia.⁴ In contrast,³ a study targeting the MDV isolation and molecular characterization in the central part of the country on infected chickens of diverse production systems grouped all isolated results in Gallid Herpesvirus type 2.

The decreasing weight, despair, crippling, bursal atrophy, neurological disease, and immediate beginning of T cell lymphomas that infiltrate lymphoid tissues, visceral organs, and peripheral nerves of infected chickens are identified as clinical signs of MD.¹¹ For instance, the chronic persistence of the virus in diseased chickens supports horizontal transmission of MDV, which advances the incidence and speeds up virus-induced tumors in multiple organs. And the connection between viral quantity or very virulent (vv) MDV and that of pathology is closely correlated.¹⁶

The World Organization for Animal Health (OIE) recommended different methods to diagnose MD, which include; indirect techniques like ELISA, AGID, and IFI. Direct diagnostic methods are; histopathology, histochemistry, PCR, virus isolation, and immuno-precipitation. More recently new techniques such as quantitative polymerase chain reaction (qPCR) and loop-mediated isothermal amplification (LAMP) which are more sensitive, specific, reliable, and faster and have been applied for the detection of MDV. Hence, molecular methods, which rapidly quantify the amount of MDV genome copies in feathers and dust are used in order to detect specific serotypes present on the broiler and layer poultry farms and for monitoring vaccination status.¹⁵

Vaccination is the extreme controlling way of Marek's disease in advancing the poultry industry. However, the major concerns are due to the latent vaccine-driven highly facilitates upsurges in pathogenicity and the achievement of controlling program by vaccination²⁸ can be challenged when there is high growing virulence of MDV strains.²³

Due to the absence of a formally organized government plan in Ethiopia, this highly contagious paralytic/oncogenic disease is causing severe economic losses resulting in serious harm to the industry of the country.⁴ The survey conducted on outbreaks of MD in a commercial poultry farm in central Ethiopia,¹⁴ investigated a 46% mortality rate, and another study⁴ on indigenous chickens in Ethiopia justified the morbidity and mortality as almost equal and as Marek's disease is still fatal even to local breeds of chickens. To control the disease by means of vaccination supported better survival. However, these vaccines are imported with high transportation costs through foreign currency, and as a result, this made the sector more expensive.¹⁶ In contrast, effective vaccines developed from local strains created great demands for genetic analysis of the virus mingling through the country.⁵

A study conducted³ in central Ethiopia, which isolated and characterized the MDV1 and recommended the isolation and molecular characterization of Marek's disease virus in other geographical areas, regarding backyard, commercial poultry farms, on different breeds and age groups. However, published information regarding isolation and molecular detection, molecular characterization, and serotype identification of MDV is scarce in southwestern Ethiopia. Thus, investigating the MD virus serotype circulating in this area is needed for good control planning and preventive measure in the country. Therefore, the objective of the current study was to perform antemortem and postmortem diagnosis, isolate, and molecular detection of Marek's disease virus in southwestern Ethiopia.

Materials and Methods

Study Areas

The present study was performed from March to November, 2021 in the Yayo from Ilubabor, Bedelle from Buno Bedelle, and Bonga town from Kaffa Zones of southwestern Ethiopia. Ilubabor and Buno Bedelle zones are found in Oromia National Regional State and While, Kaffa zone is located in Southern Nation Nationalities and People's Regional State (SNNPR). The details of study areas are explained in Table 1 as evidence obtained from their respective animal health offices.

Study Design

A case-based study was employed for this research. Before beginning any outbreak investigation, proper information regarding the outbreak was gathered from district's animal health technical personnel, officers of livestock sectors, regional laboratories, and the National Animal Health Diagnostic and Investigation Center (NAHDIC). Then, after the case-based study (outbreak investigation) was conducted with the purpose of collecting appropriate samples for isolation and molecular detection of MDV, following the outbreak information gathered.

Study Animals

The samples were collected only from the total of 50 clinically suspected chickens by MD, in an outbreak areas as 28 from Bedelle, 17 from Yayo and 5 Bonga. Three outbreaks, one from each study area were investigated. By this study, a total of 250 feathers (five from each) and six (pooled from six chickens) tissue samples of spleen, liver, and kidney; five from Bedelle and one from Yayo were collected from critically ill chickens showing clinical sign of MD. All examined chickens were exotic breeds and their age was between 12 and 13 weeks.

During the research period, the outbreak investigation was purposively employed on clinically diseased chickens in suspicion of MD. Information about the disease/outbreak was gathered by history taking from the chicken owners and collected wisely from animal health professionals working in the outbreak areas.

Clinical and Postmortem Examination of Suspected Chickens

A total of 50 sick chickens were examined for the presence of Marek's disease clinical signs such as depression, loss of weight, loss of appetite, death without showing clinical signs, paralysis of legs and wings, flaccid neck, and rising up of the skin near the hair follicles. Also, a postmortem examination was conducted on six seriously sick chickens out of 50 to observe the gross lesions. Accordingly, enlargement of visceral organs of the spleen, liver, kidney, heart, and for the presence of tumor formation in different organs.

Sample Collection and Transportation

Prior to the observation of each case, samples were collected aseptically for the aim of virus isolation, Serotype identification and molecular detection. Accordingly, spleen, and liver, kidney were collected in universal bottles and also, feather follicles of infected chickens were put into cryovials filled with viral transporting media (VTM). The collected samples were transported cold chain to the National Animal Health Diagnostics and Investigation Center (NAHDIC) laboratory. In the laboratory, they were kept at -80° C till processing, according to the world organization for animal health²⁰ terrestrial manual.

Study Site	Elevation	Latitude	Longitude	Annual Rainfall	Temperature	
					Minimum	Maximum
Yayo	1400 m	8° 26′ N	36°20°`97' E	1400 mm	18°C	27°C
Bedelle	2012 m	8°26' N	36°20' E	1361 mm	27°C	34°C
Bonga	1714 m	7°16′ N	36°14′ E	2000 mm	10.1°C	27.5°C

Table	I	Details	of	Study	Areas
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Laboratory Investigation

Cell Culture and Sample Preparation

The DF-1, passage-14 (immortalized chicken embryo fibroblast) which was gained from GD, Netherlands were recultured in 25 cm² tissue culture flask after revitalized out of liquid nitrogen. The subculture was then employed for confluent flask and tissue culture plate with six wells and preserved in Dulbecco's modified Eagle's medium (DMEM) that containing 10% fetal bovine serum at the temperature of 37°C in a humidified incubator at 5% CO_2 .

Sample Preparation

Samples collected were pooled together. The samples of tissue and feather follicle/tips broken down by sterile scalpel blade and scissor to small pieces. To facilitate maceration, the small pieces of both samples were ground by the help of mortar and pestle by sterile sand. A 10% suspension of tissue or feather tips were made in pure suspension known as phosphate buffer saline (PBS) with 2% antibiotic-antimycotic solution. The suspensions were then centrifuged at 3000 rpm for 10 min and the supernatant is harvested, filtered by 0.45 μ L Millipore filter paper and kept at -20°C prior to inoculation.²⁷

Virus Isolation

The sample suspensions prepared and stored at -20° C were thawed and 0.1 mL of the samples inoculated into confluent DF-1 cultures in six-well plates along with cell controls. After 60 min adsorption at 37°C, maintenance medium is added to each well including a negative control and incubated at 37°C in a humidified incubator at 5% CO₂. Cells were monitored every 24 h postinfection and inspected for cytopathic effects (CPEs), plaque and foci formed using an inverted microscope. On sixth day, the cultures were freeze-thawed and the resulting lysates were again inoculated into fresh cultures using 25 cm² tissue culture flasks until the third passage as this method supported the growth of MDV.^{13,18}

Virus Isolation from Feather Follicles

The feather follicle samples has been used for diagnosis of Marek's diseases virus and feather tips were used to extract the cell free MDV. Accordingly, feather tips about 5 mm were suspended in sucrose, phosphate, glutamate and albumin/ ethylenediaminetetraacetic acid (SPGA/EDTA) buffer and the buffer was added with different volumes in biosafety class-II based on laboratory procedures for the purpose of extraction and titration of cell-free virus. The suspension was filtered through a 0.45 µm membrane filter and cultured on DF-1. Then the medium was added and cultures were incubated for 15 days after 40 min absorption. This method was used to isolate serotype 1 and serotype 2 with serotype 3 (HVT) of Marek's disease virus. The CPEs that are formed as a result of MDV virus serotypes were identified properly on the basis of their foci of plaque development.²¹

Virus Isolation from Tissues

The MDV Was isolated from suspected tissue as DF-1 obtained from GD, Netherlands protocol. So, 1000 μ g/mL of streptomycin and 100 IU/mL of penicillin were added to chopped pooled tissue suspension to avoid crossing contaminants. Supernatants were harvested and inoculated on primary DF-1 cells in maintenance Dulbecco's modified Eagle's medium (DMEM) that have 2% bovine fetal calf serum (incubated at 37°C). The culture medium was observed every day until 14 days to see the visible cytopathic effects (CPE).²¹

DNA Extraction

In the NAHDIC Biotechnology laboratory, from a 10% (w/v) sample of tissue/feather, the DNA was extracted and homogenates have been carried out using the Qiagen[®] DNeasy Mini kit (cat. no. 51306, Germany) according to the manufacturing company instruction. A small piece of tissue was placed (<25 mg) in a 2 mL reaction vessel and resuspended in the tissue pellet in180 μ L in a viral lysing buffer (ATL). Then, 20 μ L of proteinase K was added and incubated at 56°C until it is lysed (1–3 h), vortices during incubation. Then, after it was centrifuged briefly below and

200 μ L buffer AL was added, vortexes for 15 seconds and incubated for 10 min at 70°C. Two hundred microliters of ethanol (96–100%) was added to the samples after brief centrifugation and then mixed for 15 seconds.

The spin column was centrifuged for one minute at 6000 (8000 rpm) and QIAamp spin was placed into clean collection tube (then collection tube with filtrate was discarded). The washing buffer (AW1) was added to opened spin column tube with volume of 500 μ L and centrifuged for one minute at 6000 (8000 rpm). Five hundred microliters of washing buffer (AW2) was added to spin column tube (located in new collection tube) following the careful opening and centrifuged for three minutes by 20,000 g (14,000 rpm). Then 200 μ L of buffer AE (elusion buffer) was added to QIAmp spin column tube and incubated for one minute at room temperature and centrifuged for one minute at 6000 (8000 rpm) and the filtrate was retained at 4°C for 48 h at the latest (prolonged storage at -20°C). At the end, the extracted DNA from isolates and fresh samples were amplified by PCR.

Real-time Polymerization Chain Reaction (RT-PCR)

The quantitative or real time PCR has been used to quantify the genome copy of MDV virus serotypes that are responsible for Marek's disease.¹ Thus, the PCR was performed targeting the MDV meq gene using forward and reverse primers of MDV1 (5'-GGAGCCGGA GAG GCTTTA TG-3') and (5'ATCTGG CCC GAATACAAG GAA-3') respectively and MDV1 Pobe 5'-(FAM) CGT CTT ACC GAG GAT CCC GAA CAG G-3'according to (Qiagen, Germany) QIAamp kit now available in NAHDIC molecular lab. All procedures were employed as recommended by kit manufacturers and the test protocol was performed with steps of initial denaturation at 95°C for 15 min followed by 45 cycles at 95°C for 15 min, annealing at 60°C for 15 min, extension at 72°C for 10 min and was cooled at 40°C.

Results

Clinical Examinations and Postmortem Findings

A total of 50 chickens (28 from Bedelle, 17 from Yayo and 5 chickens from Bonga towns) were examined (Table 2). Clinical signs of MD like; paralysis of legs and wings, gray eye, loss of weight, difficulty in breathing and depression were recorded for all the chickens sampled for this study and death of diseased chickens were reported by animal owners.

Postmortem examination was applied on six chickens with serious clinical signs of MD. Five postmortem-examined chickens were from Bedelle, one from Yayo and there were no examined chickens by postmortem from Bonga town. Upon examination, the enlargement of spleens were observed. Also, enlargement, congestion, and nodular lesion of the liver were found (Figure 1).

In general, gross Lesions observed on 3/5 from Bedelle (60%), 1/1 from Yayo (100) (Table 3) and no lesions appeared on two from Bedelle. Conversely, enlargement of spleen appeared on two chickens from Bedelle and was not observed on remaining 4 (three from Bedelle and one from Yayo).

Isolation of Marek's Disease Virus

The total of 50 pooled feather follicle samples as collected five feathers from each chicken were cultured on DF-1 cells. Out of them, MDV virus was isolated in 14/50 (28%) samples which were 9/28 (32.14%) from Bedelle town, 4/17 (23.53%) from Yayo, and 1/5 (20%) from Bonga town as shown in Table 4.

Towns	Sample Type		History of Vaccination with HVT
	Pooled Tissue	Feather Follicle	
Bedelle	5	140 (5*28)	All vaccinated
Yayo	I	85 (5*17)	Non-vaccinated
Bonga	0	25 (5*5)	All vaccinated
Total	6	250 (5*50)	256

Table 2 Number of Clinically and Postmortem Examined Chickens in the Study Areas

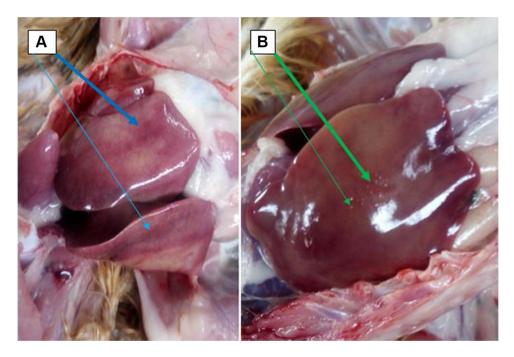


Figure I Gross lesions found on liver: (A) enlargement and congestion and (B) nodular/lymphomatous lesion.

In addition, the MDV was isolated from pooled tissue from spleen, liver, and kidney of seriously sick chickens, 5/6 chickens tissue samples from Bedelle and 1/6 tissue samples collected from Yayo. Out of six pooled tissue samples of spleen, liver, and kidney the MDV was isolated in 4/5 (80%) from Bedelle and 1/1 (100%) tissues from Yayo. In total, the virus was isolated in 5/6 (83.30%) pooled tissues of chickens (Table 5).

All of the isolated tissue samples and feather follicles showed small plaques, which are suggestive of the cytopathic effect (CPE) of MDV observed on DF-1 cells.

Initially, the CPE appeared as small and rounded cells, and at the end, it was observed as small plaque, after the formation of foci separated from the body of the culture flask. The bright and enlarged cell was also demonstrated with the help of an inverted microscope beginning from three days of the third passage as observed after 11 days of inoculation (Figure 2).

Study Site	Type of Organ Sampled	No. of Affected Organs	Type of Lesion	History of Vaccination
Bedelle	Spleen, liver, and kidney	3 liver	Lymphomatous	Vaccinated
Yayo	Spleen, liver, and kidney	I liver	Lmphomatous	Not vaccinated

Table 3 Gross Lesions on Organs of Chickens Suspected of MD from the Study Sites	Table 3 Gro	ss Lesions on	Organs of	Chickens S	Suspected (of MD	from the	Study Sites
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Study Areas	Pooled Feather Follicle	Results (CPE)	
		Number	Percent
Bedelle	28	9	32.15
Yayo	17	4	23.53
Bonga	5	I	20.00
Total	50	14	28.00

Table 4 Marek's Disease Isolation from Feather Follicle Samples

Study Areas	Pooled Feather Follicle	Results (CPE)	
		Number	Percent
Bedelle	5	4	80.00
Yayo	I	I	100.00
Total	6	5	83.30

Table 5 Marek's Disease Isolation Rate from Tissue Samples of Chicken

Real-time Polymerase Chain Reaction (RT-PCR)

A total of 50 feather follicles pooled (as collected from every 50 chickens) and six pooled tissue samples were examined by RT-PCR targeting to detect the meq gene of MDV1 using forward (5'-GGAGCCGGA GAG GCTTTA TG-3') and reverse primers of MDV1 (5'ATCTGG CCC GAATACAAG GAA-3'). DNA of 18 feather follicles was detected as MDV1 (serotype1) positive which accounts for 36%. Out of six pooled tissue samples, 5 (83.3%) of them were positive for MDV1. However, the DNA of one, 1/6 (17.3%) chicken tissue samples was not detected as MDV and resulted as negative. Thus, a summary of the PCR results was discussed in Tables 6 and 7.

All positive results of present study were indicated Gallid herpesvirus type 2 (GaHV-2), also known as Marek's disease virus 1 (Serotype-1, MDV-1). The RT-PCR results of examined samples were indicated (Figure 3).

Discussion

The clinical examination of the diseased chickens showed paralysis of legs and wings, grayness of eyes color, depression and difficulty in breathing, weight loss, and mortality. The postmortem findings in the current study also demonstrated the enlargement of the spleen, liver blood cell congestion, and lymphomatous lesion of visceral organs such as liver and lesions appeared on 4/6 of the slaughtered chickens. Both the clinical signs and postmortem examination findings indicated MDV and it was in agreement with results of previous studies.^{6,8,9,20}

In this study, MDV was isolated from pooled feather follicles of chickens which were 14/50 of sampled chickens and accounted for 28%. This finding was in agreement with previous studies,^{13,18} which isolated MDV from feather tips on the culture of DF-1. MDV was isolated from 5/6 (83.30%) of seriously ill chicken's pooled tissues of liver, spleen, and kidney. This result was related to a study for Demeke et al³ as their isolated 11/12 pooled spleen and dissimilar by types of tissues and cells. All isolates developed, small plaques, the specific and clear (CPE), which were the indicative characteristics of MDV, and this was confirmed by RT-PCR. The bright and enlarged cell was demonstrated with the help

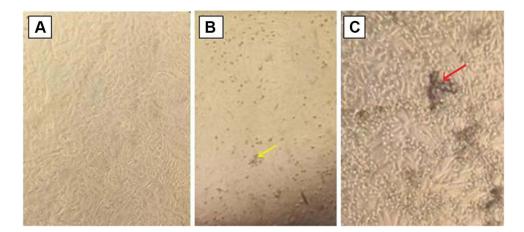


Figure 2 Non-infected and infected DFI cell by Marek's disease virus: (A) Non-infected free DF-I Cell. (B) DF-I cell infected with MDV after six days of inoculation and the yellow arrow is indicating CPE after six days and (C) CPE formed after 11 days of inoculation, cells detached and plaque formed as indicated by red arrow.

Study Areas	Feather Follicle	Results		
		No. Positive	Percent Positive	
Bedelle	28	10	35.71	
Yayo	17	7	41.18	
Bonga	5	I	20.00	
Total	50	18	36.00	

Table 6 The Summary of RT-PCR Results for Feather Follicle Samples

Table 7 The Summary of RT-PCR for Tissue Samples

Study Areas	Pooled Tissue Samples	Results		
		No. Positive	Percent Positive	
Bedelle	5	4	80.00	
Yayo	I	1	100.00	
Total	6	5	83.30	

of an inverted microscope, beginning from three days of the third passage and a clear CPE was observed after 11 days of inoculation as stated by previous studies.^{19,20,26}

The real-time PCR was used due to its high sensitivity, high specificity, efficiency, and mostly its capacity of detecting the quantity of virus accurately as it was appreciated and performed by Baigent et al.¹ By this method Marek's disease virus was detected from feather follicles of 18/50 (36%) examined chickens. This is related to the study that detected

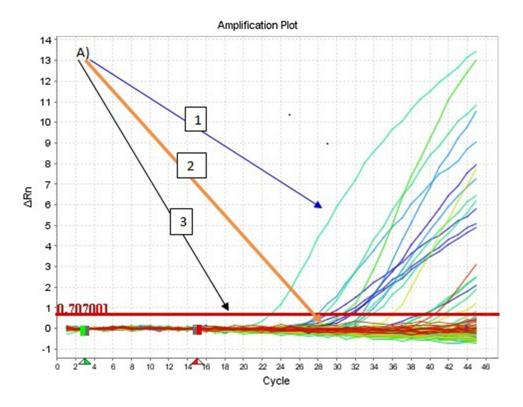


Figure 3 The real-time polymerase chain reaction results: the A1. Arrow showing the positive samples and positive control. A2. Arrow showing negative results and negative control. A3. Arrow indicating cycle threshold line.

Serotype-1 (MDV1) from feather follicles²² where less than half of the positive results were 22/173 samples tested and 13/173 tissue samples.

In addition, the Marek's disease virus was detected in pooled tissues of the chicken's spleen, liver, and kidneys as collected from critically ill chickens, during this study by RT-PCR. Accordingly, the virus was detected in 5/6 (83.30%) examined tissues of clinically suspected chickens for MD, which this result agrees with a prior study⁷ as it detected the oncogenic MDV1 virus from similar tissue samples by RT-PCR, but differs from their results which found the virus in 16/83 examined tissues.

The present study findings also confirmed the existence of Gallid herpesvirus type 2 (GaHV-2), also known as Marek's disease virus 1 (MDV-1) in southwestern Ethiopia which was devoid of previously published findings regarding MDV, based on clinical signs, isolation, and molecular detection. This result has an agreement reports^{4,14} which underlined the presence of MD in the country.

Also, the present finding was supported by that of Demeke et al,³ who conducted research on isolation and molecular characterization of MDV from clinically diseased chickens reared under different a production system in central Ethiopia and concluded that the Marek's disease virus isolates were clustered under Gallid herpesvirus type 2;¹⁶ a study based on isolation and molecular detection of MDV from chickens in the area of central Ethiopia and a study performed in central Ethiopia³ on the isolation and molecular characterization of Marek's disease which identified the presence of Meq genes of MDV1 that is circulating in the country. But, they used different diagnostic methods, specifically conventional PCR which is a qualitative method. This makes the present study findings differ as it was performed by real-time PCR.

The isolation and molecular detection of Marek's disease virus is a new one for southwestern Ethiopia as it had been justified by this study. Hence, another study focusing on viral characterization and sequencing of the strains is the key point to successful control, and prevention and better knowledge of the behavior of the virus; such as pathotypes or genotypes for eradication program.

Conclusion

The clinical and postmortem examination of the current study indicated clearly Marek's disease virus and the disease is causing morbidity and mortality of infected chickens in both vaccinated and non-vaccinated flocks in newly emerging intensive farms and backyard production systems of chickens in southwestern Ethiopia districts/towns of Bedelle, Yayo, and Bonga. Generally, the current study confirmed that the MD, which is the reason for the occurrence of an outbreak, mortality, and morbidity in chickens in the study area, was caused by Gallid herpesvirus 2 or serotype 1 (MDV-1) as it was clinically and postmortem examined, isolated and detected molecularly. Thus, this research results can contribute greatly to disease control and prevention. Hence, more extended study based on molecular characterization in the current study area, and isolation and molecular characterization of MDV in other regions of the country is recommended to identify pathotypes of the MDV that are becoming a threat to chicken production for effective control of the disease through vaccination.

Data Management and Analysis

Data obtained from laboratory tests like; virus isolation by cell culture and MDV Molecular detection by real-time polymerase chain reaction (RT-PCR) and characteristics of sampled chickens collected during sample collections were stored in Microsoft Excel prior to being analyzed using descriptive statistics and to summarize the results, the frequency table was used.

Ethical Approval Clearance

Ethical clearance was received from the National Animal Health Diagnostics and Investigation Center (NAHDIC), with reference number (ARSERC/EC/013/22/12/2020).

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Disclosure

The authors report no conflicts of interest in this work.

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