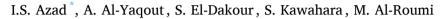
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Full Length Article First record of iridovirus (ISKNV) infections in Fourfinger threadfin from Kuwait



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ARTICLE INFO	A B S T R A C T
<i>Keywords:</i> Sheem Domestication Infection Iridovirus ISKNV	Following an initiative by the Govt. of Kuwait and the Kuwait Institute for Scientific Research (KISR), the institute started simultaneous efforts of domestication of locally caught fish and hatchery production of Four-finger threadfin (FFT, Eleutheronema tetradactylum), locally called Sheem. A private farm and KISR are working together to optimize the conditions for growth and spawning performances under the environmental conditions of Kuwait. During these efforts, infectious spleen and kidney necrosis virus (ISKNV), an important fish pathogen from Iridoviridae, caused mortalities of the FFT sub-adults (150–200 g). We noticed that the mortality was associated with anorexia, haemorrhages of the snout, fin bases, and scale pockets. We also noticed that this was typical to one of the tanks that had consistently high ammonia (0.2–0.25 ppm) levels in the water, and all 35 sub-adults died within 4–5 days. Diagnostics (molecular and electron microscopy examinations) revealed ISKN iridovirus as the causative. We noticed viral particles of 150–200 min the tissue sections of the skin, gills, heart, liver, spleen, kidney, and intestine. The major capsid protein (MCP) gene sequence analysis confirmed the results, revealing 98 % similarity with the laminin-type epidermal growth factor-like protein gene of ISKNV. The present report is the first record of ISKNV infections in FFT and from Kuwait. The study aimed to decipher the cause of illness and mortality of cultured FFT.

1. Introduction

Aquaculture, in general, is still in its infancy in the Gulf Cooperation Council member countries. Though Kuwait started aquaculture research through its Mariculture Department of KISR in the early seventies, commercial aquaculture is largely restricted to tilapia farming in the agriculture-aquaculture farming units in the Wafra region of Kuwait. The early 80s saw some efforts of cage culture in the marine environment and land-based mariculture by a private company to culture the European sea bream (*Sparus aurata*) and the local sobaity bream (*Sparidentax hasta*). However, after the Iraqi invasion, a few aquaculture entrepreneurs ventured into land-based aquaculture with Asian seabass and as a joint effort with the KISR mariculture unit with the FFT.

Fourfinger threadfin, locally known as Sheem, is a much-valued and preferred fish in the region. This fish species is available in the territorial waters of Kuwait. However, it is fast becoming scanty in Kuwait's Fishery. The Govt. of Kuwait and the KISR started an initiative for the propagation and fishery restoration efforts in 2021. Collection and domestication of the local species from Kuwait Bay began in 2021. To provide a jump-start for the research and development activity of spawning and rearing locally captured FFT, just hatched FFT larvae were procured in November 2021 from Singapore. The batch of larvae that arrived at KISR was negative for iridovirus and viral nerve necrosis through PCR diagnostics (data not shown here). The larvae were grown in land-based re-circulatory aquaculture systems (RAS). Commercial aquaculture of FFT is yet to establish itself, though scanty reports of growing this fish are available (Abu Hena et al 2011). The KISR and a private aquaculture spawning of FFT under the environmental conditions of aquaculture in Kuwait. Diseases occur in the aquaculture

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https://doi.org/10.1016/j.jksus.2024.103393

Available online 10 August 2024

Abbreviations: KISR, Kuwait Institute for Scientific Research; KFAS, Kuwait Foundation for Advancement of Science; ISKNV, Infectious Spleen and Kidney Necrosis Virus; MCP, Major Capsid Protein; PCR, Polymerase Chain Reaction; RAS, Recirculatory Aquaculture System; WOAH, World Organization for Animal Health; BHI, Brain Heart Infusion; TCBS, Thiosulphate Citrate Bile salt Sucrose; RNRS, Ribonuclease Reductase Small sub-unit; TEM, Transmission Electron Microscope; CEV, Carp Edema Virus; NCBI, National Centre for Biotechnology Information; LTEGFP, Laminin Type Epidermal Growth Factor Protein; DNA, Deoxyribose Nucleic Acid; RNA, Ribose Nucleic Acid; FFT, Fourfinger Threadfin.

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Received 20 May 2024; Received in revised form 13 July 2024; Accepted 9 August 2024

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Table 1

Primers Used for the Detection of Marine Ich and Iridovirus in Sheem.

	ase reductase small sub-unit (RNRS) NV and MCP genes of Iridovirus	Amplicon Size (bp)	References
RNRS	V1 F 5'- CACGTGTTGGCTTTCTTCGC-3'		
gene	V4 R 5'-	434	
	AGACAGGCAAAGTCACAGTG-3'	101	
	V5 R 5′ –	622	Oshima et al.,
	GAGCATCAAGCAGGCGAT-3'		1998
MCP-K1	F 5'- AAATGGCTCTTTGGAGTGTC-		Jeong et al.,
	3'		2003,2005
	R 5'- AATCCATCGGTATTATG-3'	939	

systems, especially when intensive farming practices are adopted. Viral disease such as iridovirus is very often encountered in Asian sea bass. This group of viruses is reported from fishes inhabiting different environments from freshwater to marine and from table fish production systems to aquaria (Qin et al 2023). The present investigation is aimed to

decipher the causes of illness and mortality of cultured FFT.

2. Materials and methods

Fish, Hatchlings of FFT were procured from Singapore in November 2021. The fish were reared in one-ton fiber reinforced plastic (FRP) tanks under two different systems (RAS and semi-flow-through) following optimized protocols. Temperature of the intake water, with the salinity adjusted to range between 30-32 ppt, was regulated at 27 ± 2 °C during the culture period. Briefly, the larvae were grown with green water containing nanochloropsis algal food organisms ($3x10^5$ to $5x10^5$ cells /ml) to act as food for the rotifer population ($3x10^3$ to $1x10^4$ rotifers/l). Larvae of fish were reared up to 35 days post-hatch (dph) during which time gradual weaning was carried out from 22 dph with decapsulated artemia (3-6 artemia nauplii/ml). Initial daily water exchange was 10-30 %. Feeding was done twice a day at 8.00 and 14.00 pm. Weaning to an artificial diet (BioMar, Greece) containing 42 % crude protein was initiated at 28–30 dph depending on the size groups. Rearing was continued in both production systems.

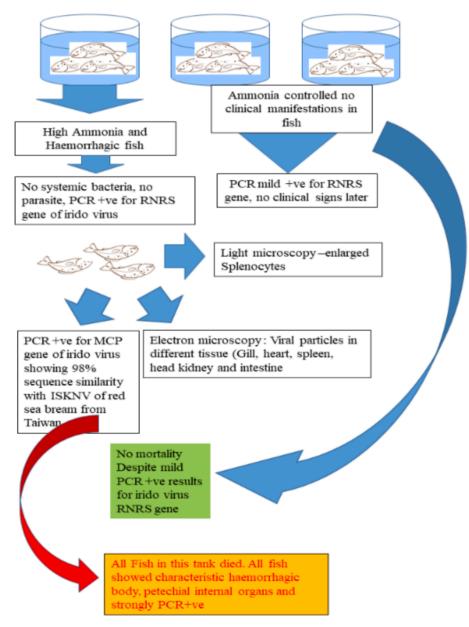


Fig. 1. Step-wise protocol of ISKNV diagnosis followed for FFT in Kuwait.

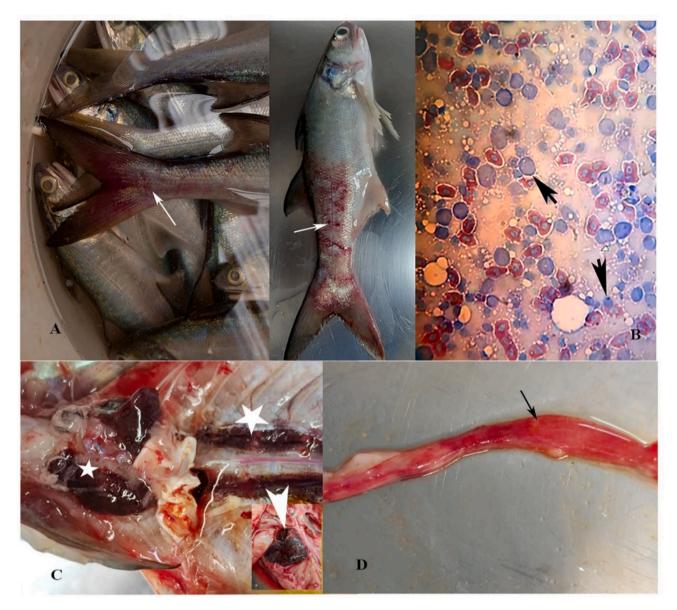


Fig. 2. Clinical manifestations in FFT with ISKNV infections. A, Gross appearance showing extensive body hemorrhage (arrows). B, Hypertrophied splenocytes (short arrows) in spleen tissue imprints (Geimsa stained). C, Enlarged head kidney (small star), necrotic mid kidney (large star), and inset showing enlarged spleen (arrowhead). D, Mid and hindgut with bloody exudates (arrow).

Clinical Manifestations, Forty-five juvenile FFT (35–43 g) were reared in each of three two-ton tanks. The tanks were fed from a storage tank, via a biofilter tank, where 100 % water exchange took place every 2–3 days. One moribund fish and two fresh-dead specimens from the tank with severe clinical manifestations were used for the investigations. Following positive diagnosis for ISKNV from the severely affected tank, two fish each from the other two tanks were also sacrificed for diagnosis.

Microbiological Observations, Aseptic blood samples from the affected fish with severe clinical manifestations were used for the detection of any possible systemic bacterial infection. Brain heart infusion (BHI) agar and thiosulphate citrate bile-salt sucrose (TCBS) agar were used for plating the samples.

Molecular Diagnosis, Samples of gills, heart, liver, spleen, kidney, and intestine were collected for both electron microscopy and for extracting DNA and PCR-based detection of the Iridovirus. Gill wet mounts were observed under the microscope to detect marine ich, and for virus detection, following primers (Table 1) were used. DNA isolation was performed using Qiagen blood/tissue DNA extraction kit following the kit protocols. Following the amplification of specific DNA

templates, the amplicons were sent to a service provider for amplicon sequencing and sequence analysis (Chromous Biotech, Bengaluru, India). Diagnostic procedures used in the present investigation were according to the Iridovirus diagnosis protocols of the World Organization of Animal Health diagnostics (WOAH, 2023) (https://www.woah. org/en/what-we-do/standards/codes-and-manuals/aquatic-manual-online-access, accessed on 8th July 2024).

Light and Electron Microscopy, Wet mounts of gills were observed under a compound microscope for recording the parasite. Spleen imprints, The fish were thoroughly bled, spleen was dissected into a sterile Petri plate, and cut into two halves one of the halves was used to get tissue imprints onto the slide, and the remaining portion was used for electron microscopy. The slide was fixed using methanol, stained with Geimsa, and observed under the microscope. The gills, heart, liver, spleen, kidney, and intestine samples were fixed in 0.3 % glutaraldehyde, critical point dried, and processed according to standard protocols for transmission electron microscopy. Thin sections were marked for observations under the transmission electron microscope (TEM). The EM facility of the Kuwait University, Health Science Centre, Jabriya,

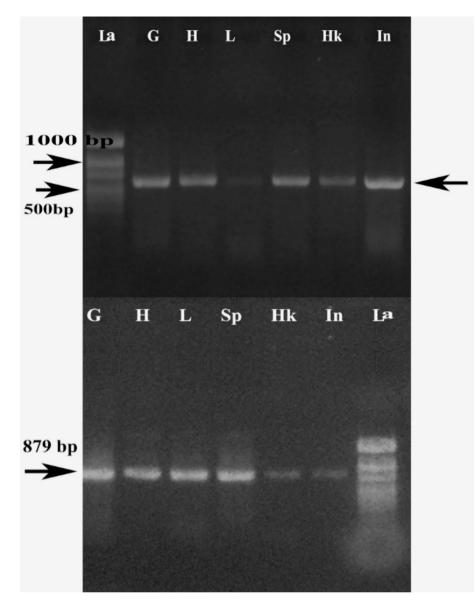


Fig. 3. PCR amplification of RNRS gene (622 bp) and ISKNV-specific MCP gene (879 bp) of infected FFT. G-gills, H-heart, L-liver, Sp –spleen, Hk-head kidney, Inintestine, La-ladder.

Kuwait was used for this purpose.

3. Results and discussions

The diagnostic procedure and infectivity confirmations were derived based on a stepwise protocol for FFT in our study (Fig. 1).

Clinical Manifestations started appearing in all three tanks in the semi flow-through system, which recorded elevated ammonia levels of 0.15- to 0.2 ppm. The fish showed reduced appetite, irritability, and reddening. The storage tank water was completely replenished following the symptoms. Ammonia levels returned to less than 0.1 ppm following replenishment. Despite these efforts, one tank continued to have elevated levels of ammonia (>0.1 ppm). Clinical symptoms in this tank became more intense with hemorrhagic snout, fin bases, and distal body surface (Fig. 2). While. clinical manifestations of the fish in the other two tanks became normal as the ammonia levels were consistently lower than 0.1 ppm. All the fish in the severely affected tank died within 4–5 days despite efforts to keep the ammonia levels low and preventive measures (oxytetracycline bath 40 ppm daily) to control opportunistic vibriosis.

Hemorrhagic body surface, fin bases, and snout were the consistent clinical manifestations in all the fish. Internal organs showed petechial hemorrhages, pale liver, enlarged spleen, and empty gut with the hemorrhagic midgut. The spleen tissue imprint showed basophilic inclusion bodies (Fig. 2). Clinical manifestations in FFT, as observed in the present case, match the manifestations reported for several fish species (Kurita, 2012: Subramaniam et al., 2012: Dong et al., 2017: OIE, 2019). Spleen and kidney enlargement seen in FFT are similar to the observations made by He and his team (He et al., 2000 and 2002). Mandarin fish and several other marine fish, under an experimental infection model, showed spleen, kidney, and endocardial hypertrophy as common ISKNV-associated manifestations. Total heterotrophic bacterial counts in the affected tank water were $1.2 \pm 6.3 \times 10^3$ CFU/ml and the total vibrio counts were < 100 CFU/ml. Neither systemic bacterial infection nor parasitic infestation was recorded from the infected fish. There was no significant difference between the bacterial counts of different tanks. In our case study, we see a relation between the elevated water ammonia levels and the outbreak of ISKNV. However, in the tanks where ammonia levels triggered clinical manifestations and the virus was detected in the FFT, improvement in the water quality was probably responsible for

	Large yellow croaker iridovirus isolate ZS-2, complete genome
	viruses 9 leaves
	Red sea bream iridovirus genomic DNA, circular physical map, complete sequence
	Large yellow croaker iridovirus, complete genome
	Red sea bream iridovirus strain Namhae unknown genes
_	viruses 11 leaves
	Megalocytivirus FD201807, complete genome
	viruses 4 leaves
	Banggai cardinalfish iridovirus isolate BCIV/WVL17393/2012
	Infectious spleen and kidney necrosis virus isolate BCIV-2015, complete genome
	Largemouth bass ulcerative syndrome virus isolate LBUSV-GZ, complete genome
Ť	Infectious spleen and kidney necrosis virus isolate OGIV-YK-2018, complete genome
	Infectious spleen and kidney necrosis virus isolate HGIV-2015, complete genome
	Infectious spleen and kidney necrosis virus isolate XMIV-2015, complete genome
	Infectious spleen and kidney necrosis virus isolate TLIV-2015, complete genome
	Infectious spleen and kidney necrosis virus isolate XHIV-2015, complete genome
	Infectious spleen and kidney necrosis virus isolate DGIV_2010, complete genome
	Infectious spleen and kidney necrosis virus isolate NH-2005, complete genome
	Infectious spleen and kidney necrosis virus isolate TTIV-2011, complete genome
	Infectious spleen and kidney necrosis virus, complete genome
	Infectious spleen and kidney necrosis virus strain RSIV-Ku
	Infectious spleen and kidney necrosis virus strain TIV-2020, complete genome
	Spotted knifejaw iridovirus isolate SKIV-SD
	Infectious spleen and kidney necrosis virus isolate Bali/Hybrid-grouper/2016/SVC-18-009
	Infectious spleen and kidney necrosis virus isolate ISKNV_Ghana_TIV_2019, complete genome
	PInfectious spleen and kidney necrosis virus strain LakeVolta_BF-2, complete genome
	Infectious spleen and kidney necrosis virus isolate KU2
	Infectious spleen and kidney necrosis virus isolate KU1
	Infectious spleen and kidney necrosis virus isolate EFIV-2019
	Infectious spleen and kidney necrosis virus isolate NH-2021, complete genome
	PAngelfish iridovirus AFIV-16
	Infectious spleen and kidney necrosis virus isolate OGIV-PW-2018, complete genome
	Infectious spleen and kidney necrosis virus isolate TTIV-2015, complete genome
	Infectious spleen and kidney necrosis virus strain AK-ISKNV, complete genome
	Infectious spleen and kidney necrosis virus isolate HGIV-Cantik1-2014, complete genome
	Infectious spleen and kidney necrosis virus isolate TLIV-2011, complete genome
	Infectious spleen and kidney necrosis virus isolate EFIV-2018
	Infectious spleen and kidney necrosis virus, complete genome
	♥Infectious spleen and kidney necrosis virus isolate Bali/ Hybrid-grouper/2016/SVC-18-072 ♥Banggai cardinalfish iridovirus isolate BCIV/2017
0.01	
	Infectious spleen and kidney necrosis virus isolate KISR_ISKNV1 laminin-type epidermal growth factor-like protein gene, partial cds

Fig. 4. Distance tree of ISKNV of FFT from Kuwait.

preventing the seriousness of the infection, and no mortality was recorded from those tanks. It is well known that the environment influences the pathogen invasiveness (Engering et al., 2013). Such an environment-triggering effect was demonstrated in an experiment where dissolved oxygen levels influenced ISKNV infectivity (Yu et al., 2022). Exposure to stress and endogenous ammonia adversely affects the ability of the fish to maintain homeostasis (Randall and Tsui, 2002). They observed that apart from the metabolic stressors, hyper-ammonia condition was a predisposing factor in the carp edema virus (CEV) infections in carp. Similarly, metabolic disturbances were also reported to be one of the pre-disposing factors in CEV (Pikula et al., 2021).

Fourfinger threadfin is gaining importance as a food fish following its controlled seed production and larval rearing efforts (World Aquaculture Society, 2024). Aquaculture production of FFT in Kuwait is in its early days thanks to the Government initiative. This species is known to be a stress-sensitive fish as we noticed ammonia-mediated necrotic lesions in this fish during the early development period. Similar stress-induced immune compromises have been reported from Atlantic salmon (Tamsyn et al, 2021). Molecular diagnostics using primers from published and related work (Oshima et al. 1998) produced positive amplification for the iridovirus (Fig. 3). The sequence analysis of the 622 bp amplicon produced 99 % sequence similarity with the angelfish iridovirus RNRS gene (MK689685.1). Hence another set of primers (Jeong et al., 2003) targeting the major capsid protein gene of ISKNV was used for specific diagnostics. These reactions produced 879 bp amplicon (Fig. 2) and a 98 % sequence similarity with MCP gene sequence of red seabream ISKNV from Taiwan (KT781098.1). The 879 bp MCP gene sequence of the ISKNV in FFT from Kuwait was submitted to the NCBI (OR136938.1).

The distance tree is depicted in Fig. 4. The gene sequence corresponded to the gene sequence of the laminin-type epidermal growth factor-like protein (LTEGFP). The MCP of the megalocytivirus, to which the ISKNV belongs, contains 40 % of the proteins with more than 36 additional polypeptides. These polypeptides are known to take part in viral particle formation (Robin et al., 1986). The MCP gene sequence of ISKNV of FFT, corresponding to the LTEGFP gene sequence, indicates a possible role of this protein in the infectivity of the virus. The involvement of the LTEGFP in providing a mock basement membrane that

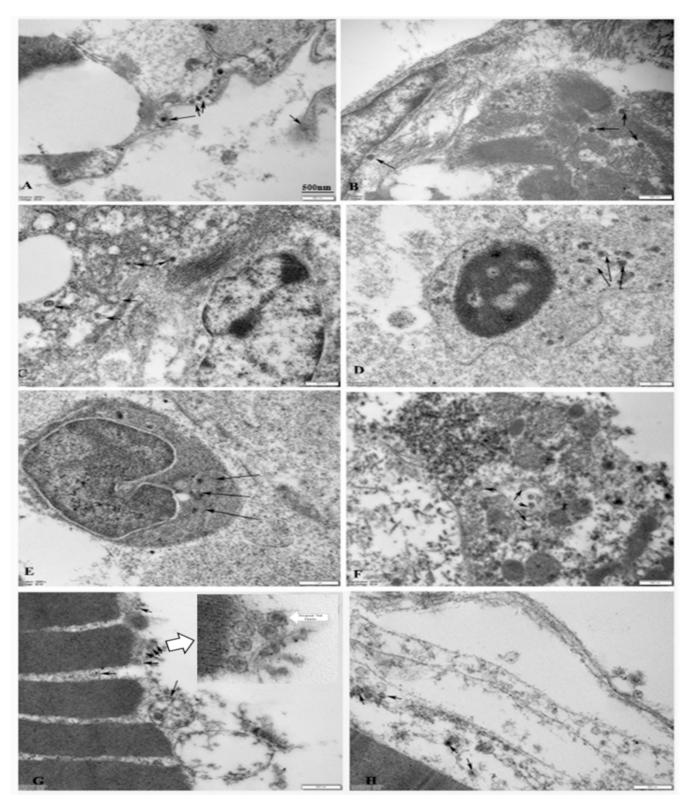


Fig. 5. Viral infections as seen under the transmission electron microscope. A-Gill epithelium containing viral particles (arrows). B –Myocardium showing viral particles in the myofiber interstitium (arrow). C –Liver tissue with viral particles (arrows). D –Splenocyte showing intracytoplasmic localization of viral particles (arrows). E –pronephric lymphopoietic tissue with viral inclusion (arrows). F- Intestinal epithelium with viral particles (arrows). G –Epidermis and basement membrane with viral particles (arrows).

facilitates viral attachment to lymphatic endothelial cells has been demonstrated earlier (Xu et al 2010).

Electron Microscopy of different tissues of FFT (Fig. 5) revealed the presence of viral particles associated primarily with the epithelial cells

of gills, myocardial interstitial spaces, hepatic acini, splenocytes, pronephric lymphoid tissue, intestinal epithelium, and muscle-epidermal basement. The viral particles were associated with the inclusion bodies within various cells of the gills, heart, liver, spleen, kidney, intestine, and muscle. It will be interesting to see if this LTEGPF gene corresponds to the VP23R protein (Xu et al., 2010) reported to have an ISKNV infection-related role in ISKNV infectivity. There are no reports of ISKNV from cultured FFT. However, this species has been listed as susceptible to ISKNV by the WOAH in its report (WOAH, 2023). The first report of FFT susceptibility to viral infections from cultured fish indicated RNA viruses cause pathological manifestations in FFT from Singapore (Chang et al., 2002). The list of species susceptible to ISKNV (WOAH, 2023) reflects the possibility of infections in natural environments and culture conditions. The carrier state of the cultured organisms, either through the environment or via parental transfer, is one of the potential sources of the disease. Limitations of detection specificity and sensitivity of ISKNV, in stockable fish, could also contribute to the disease situation when culture conditions become stressful to fish. The possible role of the integument and the gill epithelium in becoming portals of entry for the ISKNV cannot be ruled out as the viral particles were seen associated with the basement membranes of the skin and epithelial lining of the gills in our study.

4. Conclusion

Fourfinger threadfin is sensitive to handling and water quality parameters. The aim and the purpose of the present investigation of deciphering the cause of illness and mortality were achieved. A greater purpose was also met by sequencing the major capsid protein of the virus for future work on vaccine development. Infectivity and resultant moralities could occur due to ISKNV in this fish. Hence, water quality management and preventive vaccination procedures must be applied in commercial aquaculture of this species.

CRediT authorship contribution statement

I.S. Azad: Validation, Supervision, Methodology, Investigation, Data curation, Conceptualization. A. Al-Yaqout: Supervision, Methodology, Funding acquisition. S. El-Dakour: Supervision, Methodology. S. Kawahara: Validation, Supervision, Investigation. M. Al-Roumi: Supervision, Resources.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Data availability

Data will be made available on request.

Acknowledgment

We thank Ms. Jessy and the Head of the Electron Microscopy Unit of Kuwait University, Health Science Centre, Jabriya, Kuwait for extending their facilities for microscopic investigations. The authors are thankful to Kuwait Foundation for Advancement of Science (KFAS) for partial funding support.

Appendix A. Supplementary material

Supplementary data to this article can be found online at https://doi.org/10.1016/j.jksus.2024.103393.

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Further Reading

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