Conference on Veterinary Sciences

MDPI

03-05 December 2025 | Online

Molecular Characterisation of Fowl Adenoviruses Associated with Inclusion Body Hepatitis and Gizzard Erosion in Broiler Chickens in Sabah, Malaysia

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INTRODUCTION & AIM

- Inclusion Body Hepatitis (IBH) and adenoviral gizzard erosion (AGE) are major viral diseases with significant health concerns in poultry production worldwide (Sohaimi & Ugwu, 2021).
- Both conditions are linked to fowl adenoviruses (FAdVs), but serotypes circulating in Sabah remain poorly characterised.

AIM: To identify and characterise FAdV serotypes associated with IBH and AGE in broiler chickens between 2019 and 2023.

METHOD

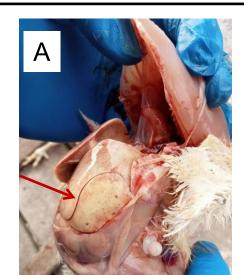


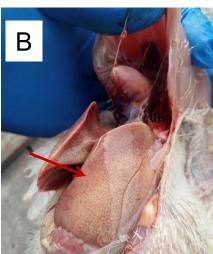
FIGURE 1: Map of Sabah, Malaysian Borneo and location of sample collection throughout the state .

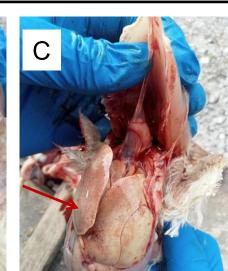
- A total of 66 liver and gizzard samples collected from suspected IBH or AGE cases (2019–2023).
- PCR screening for FAdV using hexon-targeting primers (Meulemans et al., 2001).
- Samples underwent partial sequencing of the hexon gene, a major capsid protein used for FAdV serotyping.
- Sequence analysis was performed to detect nonsynonymous amino acid variants within serotypes.

RESULTS & DISCUSSION

- 49 of 66 samples tested positive for FAdV.
- Three serotypes identified: FAdV-8b, FAdV-1, FAdV-11.

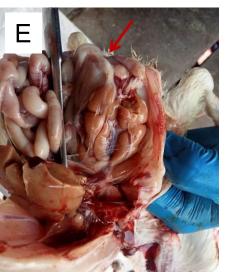








A-C: Liver Hepatitis



F





G: Breast and peritoneal muscles hemorrhage

H: Thigh muscle hemorrhage

Yellowish

Diarrhea



E: Swollen Kidney F: Jaundice

I: Subcutaneous Jaundice





FIGURE 2: Morphological features and physical symptoms of IBH and AGE Includes A) Liver Hepatitis, B) Jaundice, C) Gizzard Erosion, D) Swollen Kidney, F) Haemorrhage of the Breast, Thigh and Peritoneal Muscle, G) Yellowish Diarrhoea, H) Weakness and Depression

J: Weak and depressed birds

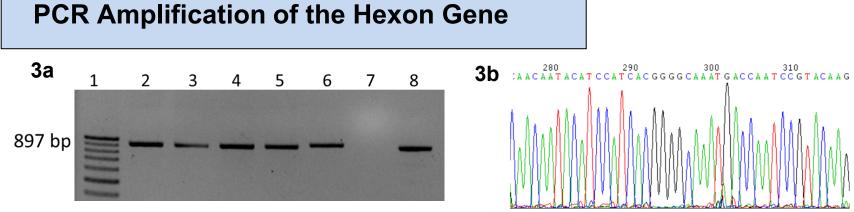
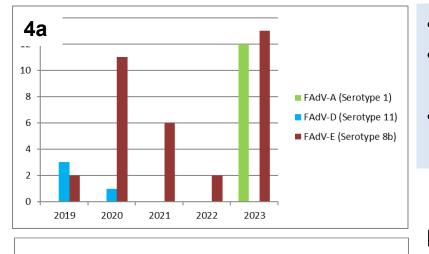


FIGURE 3: a) PCR amplification of the Hexon gene showing an expected 897 bp amplicon. Lane 1: 100 bp DNA ladder; Lanes 2–6: liver tissue samples; Lane 7: negative control; Lane 8: positive control. **b)** DNA sequence chromatogram of the Hexon gene.

Distribution of FAdV Serotypes Detected in Sabah (2019–2023)



- FAdV-11 restricted to 2019–2020.
 FAdV-1 detected only in 2023, coinciding with
- AGE-like gizzard lesions.
 Surge in IBH cases in 2023 aligned with increased FAdV-8b detection.

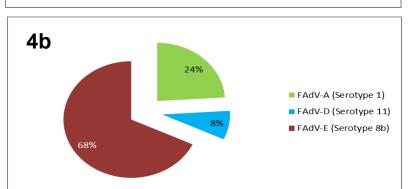


FIGURE 4: a) Annual distribution of detected Fowl adenovirus serotypes from 2019 to 2023 show distinct temporal patterns, with FAdV-E predominating across multiple years. **b)** Overall proportion of FAdV serotypes detected across all sampled years.

ML tree of the 49 Field Isolates Mapped to Reference Strains.

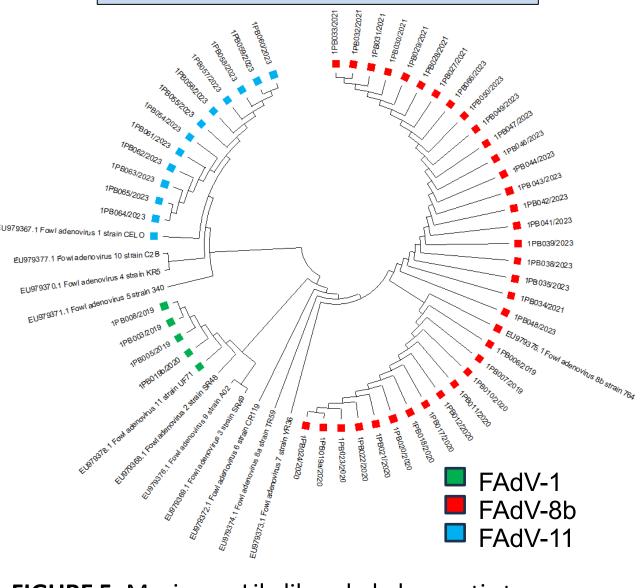


FIGURE 5: Maximum Likelihood phylogenetic tree (using the Tamura-Nei model) clearly delineates the 49 samples into FadV-1, FAdV-8b and FADV-11 serotypes.

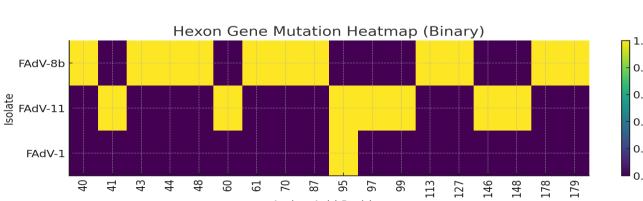


Table 1: Hexon Amino-Acid Substitutions in Sabah FAdVs

Serotype	Non-Synonymous
	Amino Acid changes
FAdV-8b	140V, D43E, G44D, T48S,
	A61Q, A70G, G87A, N113S,
	L127I, G178R, E179A
FAdV-11	E41D, N60V, E95A, T97S,
	V99A, G146A, K148N
FAdV-1	A95P

- Multiple N-terminal changes in 8b and 11A & distinct A95P substitution in FAdV-1.
- Sabah isolates indicate antigenic variation, particularly within the L1 region, which may influence immune recognition.
- Hexon mutations may indicate:
- a. Antigenic drift / immune escape
- b. Possible changes in host immune response
- c. Serotype-level diversity and evolutionary pressure (Niczyporuk, 2018).

FIGURE 6: Amino acid heatmap indicating location of Mutations in the Sabah isolates of the FAdV-1, FAdV-11 and FAdV-8b stains (Yellow = mutations; Purple = Reference strains- Wildtype).

CONCLUSION

- First molecular evidence of circulating FAdV-1, -8b, and -11 in Sabah broiler flocks.
- Serotype prevalence shifted over the 5-year period, with FAdV-8b becoming dominant.
- Mutation Hexon mutations observed in Sabah isolates indicate antigenic variation, which may influence immune recognition.
- Findings highlight the need for routine molecular surveillance and improved farm biosecurity (Kardoudi *et al.*, 2025).

REFERENCES

ACKNOWLEDGEMENTS

- Kardoudi *et al.*, 2025. Veterinary Sciences12(4):378. We thank the various poultry farms for Meulemans *et al.*2001. Avian Pathology 30(6):655-60. providing samples. The project was
- Niczyporuk JS. 2018. PLoS One 13(11):e0207668. Sohaimi & Ugwu. 2021. Journal of World's Poultry Research 11(3):387-396.
- providing samples. The project was funded by internal grants from the Biotechnology Research Institute, UMS.