



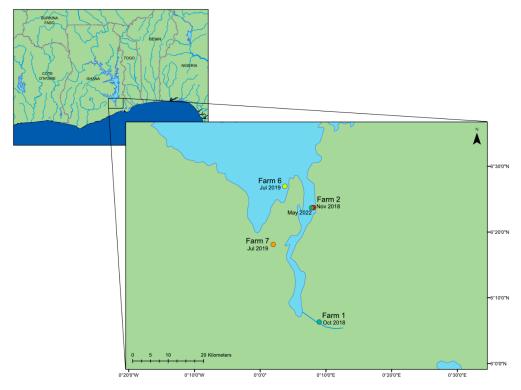
Correction

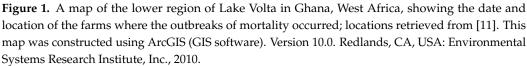
Correction: Alathari et al. A Multiplexed, Tiled PCR Method for Rapid Whole-Genome Sequencing of Infectious Spleen and Kidney Necrosis Virus (ISKNV) in Tilapia. *Viruses* 2023, 15, 965

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In the original publication [1], there was a mistake in Figures 1 and 2 as published. In Figure 1, there is a loss of colour in most of the dots representing the farms on the map. The corrected Figure 1 appears below.





In Figure 2, there is a loss of all dots. The corrected Figure 2 appears below.



Citation: Alathari, S.; Chaput, D.L.; Bolaños, L.M.; Joseph, A.; Jackson, V.L.N.; Verner-Jeffreys, D.; Paley, R.; Tyler, C.R.; Temperton, B. Correction: Alathari et al. A Multiplexed, Tiled PCR Method for Rapid Whole-Genome Sequencing of Infectious Spleen and Kidney Necrosis Virus (ISKNV) in Tilapia. Viruses 2023, 15, 965. Viruses 2023, 15, 1476. https://doi.org/10.3390/v15071476

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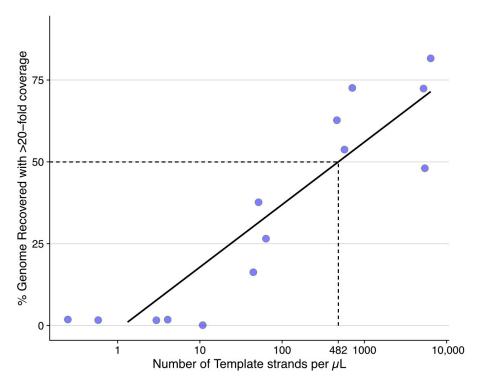


Figure 2. Successful recovery of >50% of the ISKNV genome required 482 template strands per μ L (2410 viral templates per 5 μ L sequencing reaction), with a minimum of 0.2 copies per μ L to recover >0% of the genome with at least 20-fold coverage for error correction. Number of viral templates was measured using ddPCR from a serially diluted ISKNV template, which was subsequently sequenced and processed as described in the text.

The authors state that the scientific conclusions are unaffected. This correction was approved by the Academic Editor. The original publication has also been updated.

Reference

1. Alathari, S.; Chaput, D.L.; Bolaños, L.M.; Joseph, A.; Jackson, V.L.N.; Verner-Jeffreys, D.; Paley, R.; Tyler, C.R.; Temperton, B. A Multiplexed, Tiled PCR Method for Rapid Whole-Genome Sequencing of Infectious Spleen and Kidney Necrosis Virus (ISKNV) in Tilapia. *Viruses* 2023, 15, 965. [CrossRef] [PubMed]

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