



# Correction for Batovska et al., “Coding-Complete Genome Sequence of Yada Yada Virus, a Novel Alphavirus Detected in Australian Mosquitoes”

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Volume 9, no. 2, e01476-19, 2020, <https://doi.org/10.1128/MRA.01476-19>. After publication, the authors were made aware of a short sequence on GenBank ([JQ749729.1](https://www.ncbi.nlm.nih.gov/nuclseq/JQ749729.1)) with high similarity to Yada Yada virus (YYV). The 331-bp sequence is part of the nonstructural polyprotein (NSP4) and shares 97% nucleotide identity and 99.1% amino acid identity with YYV. This sequence was detected in an *Aedes notoscriptus* mosquito collected in North Melbourne, Australia, in 2011. The similarity of this sequence to YYV suggests that the virus from which it originated is part of the mosquito-specific alphavirus complex, which consists of viruses that have been detected only in *Anopheles* and *Culex* mosquitoes.

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