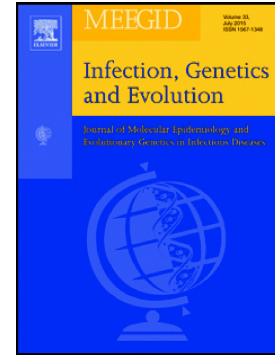


Accepted Manuscript

High-resolution phylogeny providing insights towards the epidemiology, zoonotic aspects and taxonomy of sapoviruses

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PII: S1567-1348(17)30330-1
DOI: doi:[10.1016/j.meegid.2017.09.024](https://doi.org/10.1016/j.meegid.2017.09.024)
Reference: MEEGID 3280
To appear in: *Infection, Genetics and Evolution*
Received date: 23 August 2017
Revised date: 18 September 2017
Accepted date: 19 September 2017

Please cite this article as: A.F. Barry, R. Durães-Carvalho, E.F. Oliveira-Filho, A.A. Alfieri, W.H.M. Van der Poel , High-resolution phylogeny providing insights towards the epidemiology, zoonotic aspects and taxonomy of sapoviruses. The address for the corresponding author was captured as affiliation for all authors. Please check if appropriate. Meegid(2017), doi:[10.1016/j.meegid.2017.09.024](https://doi.org/10.1016/j.meegid.2017.09.024)

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**High-resolution phylogeny providing insights towards the epidemiology, zoonotic aspects
and taxonomy of sapoviruses**

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Abstract

The evolution, epidemiology and zoonotic aspects of Sapoviruses (SaV) are still not well explored. In this study, we applied high-resolution phylogeny to investigate the epidemiological and zoonotic origins as well as taxonomic aspects of animal and human SaV. Bayesian framework analyses showed an increase in porcine SaV (PoSaV) population dynamics between

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