

Correction

Correction: Zablocki, O.; *et al.* Diversity of dsDNA Viruses in a South African Hot Spring Assessed by Metagenomics and Microscopy. *Viruses* 2017, 9, 348

Olivier Zablocki , Leonardo Joaquim van Zyl, Bronwyn Kirby and Marla Trindade * 

Institute for Microbial Biotechnology and Metagenomics, Department of Biotechnology, University of the Western Cape, 7535 Bellville, South Africa; olivier.zablo@gmail.com (O.Z.); vanzylj@gmail.com (L.J.v.Z.); bkirby@uwc.ac.za (B.K.)

* Correspondence: ituffin@uwc.ac.za; Tel.: +27-021-959-9725

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The authors wish to make the following changes to their paper [1]. In Section 3.5.1. (Identification Using the *polB2* Gene), we erroneously grouped His1 and His2 viruses in the same genus, *Salterprovirus*, and had concluded that our metagenomic sequences which had grouped with His1 and His2 viruses had expanded the *Salterprovirus* clade (Figure 2, [1]). His1 virus is the sole current representative of the genus *Salterprovirus* [2]. However, His2 virus is classified as belonging to the genus *Gammapleolipovirus* [3]. Thus, metagenomic sequences related to both His1 and His2 viruses may be expanding either genera or both, and not only the genus *Salterprovirus*, as initially stated in the submitted paper.

The authors apologize for any inconvenience this may have caused. The changes do not significantly affect the scientific results however rectify the taxonomic inaccuracies.

References

1. Zablocki, O.; van Zyl, L.J.; Kirby, B.; Trindade, M. Diversity of dsDNA Viruses in a South African Hot Spring Assessed by Metagenomics and Microscopy. *Viruses* **2017**, *9*, 348. [[CrossRef](#)] [[PubMed](#)]
2. Hong, C.; Pietilä, M.K.; Fu, C.J.; Schmid, M.F.; Bamford, D.H.; Chiu, W. Lemon-shaped halo archaeal virus His1 with uniform tail but variable capsid structure. *Proc. Natl. Acad. Sci. USA* **2015**, *112*, 2449–2454. [[CrossRef](#)] [[PubMed](#)]
3. Pietilä, M.K.; Roine, E.; Sencilo, A.; Bamford, D.H.; Oksanen, H.M. Pleolipoviridae, a newly proposed family comprising archaeal pleomorphic viruses with single-stranded or double-stranded DNA genomes. *Arch. Virol.* **2016**, *161*, 249–256. [[CrossRef](#)] [[PubMed](#)]



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