

**Part 1:** **TITLE, AUTHORS, APPROVALS, etc**

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| **Code assigned:** | **2021.069B** |  |
| **Short title:**  Rename the genus *Giessenvirus* (to *Hungariovirus*) (*Caudoviricetes*) | | |
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**Author(s) and email address(es)**

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| Sváb D, Tóth I | svab.domonkos@atk.hu; tothistvan3@yahoo.co.uk |

**Author(s) institutional address(es) (optional)**

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| Institute of Veterinary Medical Research, Centre for Agricultural Research, Budapest, Hungary [SD, TI] |

**Corresponding author**

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| Domonkos Sváb |

**List the ICTV Study Group(s) that have seen this proposal**

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| Bacterial Viruses Subcommittee |

**ICTV study group comments and response of proposer**

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**Authority to use the name of a living person**

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| **Is any taxon name used here derived from that of a living person (Y/N)** | N |

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| **Taxon name** | **Person from whom the name is derived** | **Permission attached (Y/N)** |
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**Submission dates**

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| Date first submitted to SC Chair | March 30, 2021 |
| Date of this revision (if different to above) |  |

**ICTV-EC comments and response of the proposer**

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| Acceptance of proposal 2021.001B.abolish\_Caudovirales by EC53 results in removal of the order *Caudovirales* and families *Myoviridae*, *Podoviridae* and *Siphoviridae*. All underlying taxa are to be assigned directly to the class *Caudoviricetes*. The Excel module of this proposal has been altered to reflect the future changes; however, the Word module has been unaltered while awaiting the ratification vote. |

**Part 3:** **TAXONOMIC PROPOSAL**

**Name of accompanying Excel module**

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| 2021.069B.R.Rename\_Giessenvirus |

**Abstract**

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| We propose the renaming of the genus *Giessenvirus* to *Hungariovirus*, as the earlier name was proposed based on the supposed place of isolation of the representative bacteriophage isolate (C130\_2), which, contrary to the assumption of the original proposal, was Budapest, Hungary, therefore a new genus name is proposed. |

**Text of proposal**

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| |  | | --- | | We ask for the renaming of the genus ‘*Giessenvirus*’, established in proposal no. 2019.076B. [1], with its sole representative phage C130\_2, which was described by Sváb et al., in Arch. Virol., 2019 [2].  In the proposal it is stated that phage C130\_2 was isolated in Giessen, Germany. This is not correct, and it is supposedly due to the fact that in the original paper we did not specify where each part of the work was performed. DNA sequence determination and the associated bioinformatics work, including uploading the sequence to GenBank was indeed performed at the University of Giessen. However, isolation, purification and phenotypic characterization of the phage, which included the determination of host spectrum, was performed at the Institute for Veterinary Medical Research, Centre for Agricultural Research, Budapest, Hungary. Therefore, we propose the genus name, *‘Hungariovirus*’ for bacteriophage C130\_2. | |

**Supporting evidence**

**References**

1. Kropinski AM, Adriaenssens EM (2019) Create four new genera including six new species in the family Podoviridae 2019.076B. Phage.Canada@gmail.com. [https://ictv.global/ICTV/proposals/2019.076B.zip. Accessed 30 March 2021](https://ictv.global/ICTV/proposals/2019.076B.zip.%20Accessed%2030%20March%202021)
2. Sváb D, Falgenhauer L, Rohde M, et al (2019) Complete genome sequence of C130\_2, a novel myovirus infecting pathogenic *Escherichia coli* and *Shigella* strains. Arch Virol 164:321–324. <https://doi.org/10.1007/s00705-018-4042-0>. PMID: 30238162