

The International Committee on Taxonomy of Viruses

Taxonomy Proposal Form, 2024

**Part 1a: Details of taxonomy proposals**

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| **Title:**  | Edit a typo in all the 19 species epithets of subfamily *Spumaretrovirinae*, specifically one species in genus *Bovispumavirus*, *Equispumavirus*, *Felispumavirus*, *Prosimiispumavirus* each, and 15 species in genus Simiispumavirus. |
| **Code assigned:**  | 2023.009DX.Retroviridae\_68rensp\_Error\_Correction |

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| **Author(s), affiliation and email address(es):**  |
| **Name**  | **Affiliation**  | **Email address**  | **Corresponding author(s)** X |
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**Part 1b: Taxonomy Proposal Submission**

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| **ICTV Subcommittee:**  |
| Animal DNA Viruses and Retroviruses | **x** | Bacterial viruses |  |
| Animal minus-strand and dsRNA viruses |  | Fungal and protist viruses |  |
| Animal positive-strand RNA viruses |  | Plant viruses |  |
| Archaeal viruses |  | General - |  |

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| **List the ICTV Study Group(s) that have seen or have been involved in creating this proposal:**  |
| *Retroviridae* SG |

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| **Optional – complete only if formally voted on by an ICTV Study Group:**  |
| **Study Group** | **Number of members** |
| **Votes in support** | **Votes against** | **No vote** |
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| **Submission date:** |  13/05/2024 |

**Part 1c: Feedback from ICTV Executive Committee (EC) meeting**

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| **Executive Committee Meeting Decision code:** | **X** |
| A – Accept |  |
| Ac – Accept subject to revision by relevant subcommittee chair. No further vote required |  |
| U – Accept without revision but with re-evaluation and email vote by the EC |  |
| Uc – Accept subject to revision and re-evaluation and email vote by the EC |  |
| Ud – Deferred to the next EC meeting, with an invitation to revise based on EC comments |  |
| J - Reject |  |
| W - Withdrawn |  |

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| **Comments from the Executive Committee:** |
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**Part 1d: Revised Taxonomy Proposal Submission**

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| **Response of proposer:**  |
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| **Revision date:** |  DD/MM/YYYY |

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

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| **Name of accompanying Excel module:**  |
| 2023.009DX.Retroviridae\_68rensp\_Error\_Correction.xlsx |

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| **Taxonomic changes proposed:**  |
| Establish new taxon |  | Split taxon |  |
| Abolish taxon |  | Merge taxon |  |
| Move taxon |  | Promote taxon |  |
| Rename taxon | **x** | Demote taxon |  |
| Move and rename |  |

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| **Is any taxon name used here derived from that of a living person:**  |  **N** |
| **Taxon name** | **Person from whom the name is derived** | **Attached X** |
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| **Abstract of Taxonomy Proposal:**  |
| *Taxonomic rank(s) affected*: Species*Description of current taxonomy*: 19 species classified in 5 genera within one subfamily.*Proposed* *taxonomic change(s):* Error correction of a typo on 19 species epithets*Justification*:An error has come across in the 2023.009D.A.v2.Retroviridae\_68rensp.xlsx validation, with the epithet of all 19 species in subfamily Spumaretrovirinae erroneously ending in "…fo". We would like to correct this, that is, delete "fo" throughout, in order to reflect the correct species names.  |

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| * **Text of Taxonomy proposal:**
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| *Taxonomic rank(s) affected*: Species*Description of current taxonomy*: 19 species classified in 5 genera within one subfamily.*Proposed* *taxonomic change(s)*: Edit 19 species names to lack "fo" as the last two letters in the specie epithet.*Demarcation criteria:**n/a**Justification*: An error has come across in the 2023.009D.A.v2.Retroviridae\_68rensp.xlsx validation, with the epithet of all 19 species in subfamily Spumaretrovirinae erroneously ending in "…fo". We would like to correct this, that is, delete "fo" throughout, in order to reflect the correct species names.  |

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| **References:**  |
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| **Tables, Figures:**  |

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