

The International Committee on Taxonomy of Viruses

Taxonomy Proposal Form, 2025

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

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| **Title:**  | Create one new species in the genus *Deltasatellite* (*Tolecusatellitidae)* |
| **Code assigned:**  | 2025.015P.Ac.v3.Deltasatellite\_1nsp |

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| **Author(s), affiliation and email address(es):**  |
| **Given name (+middle initial(s))** | **Surname** | **Affiliation**  | **Email address**  | **Corr. author(s)**  |
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**Part 1b: Taxonomy Proposal Submission**

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

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

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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** |
| Roumagnac, PhilippeAscencio-Ibáñez, Jose TLett, Jean-MichelLópez-Lambertini, Paola M.Martin, DarrenNavas-Castillo, JesúsRibeiro, SimoneUrbino, CicaVarsani, ArvindZerbini, F. Murilo | YYYYYYYYYY |  |  |

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| **Submission date:** |  05/06/2025 |

**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 | **X** |
| 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| W - Withdrawn |  |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:** |
| The EC voted Ac for this proposal (see the table above for explanation), to allow very minor revisions mainly concerning style issues. |

**Part 1d: Revised Taxonomy Proposal Submission**

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| **Response of proposer:**  |
| All style-related suggestions were accepted.  |

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| **Revision date:** | 22/08/2025 |

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

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

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| **Etymology (origin) of proposed taxonomic names:**  |
| **Taxon name**  | **Etymology of the term** |
| *“Deltasatellite brassicae”* | The species epithet (*brassicae*) refers to the begomovirus species to which it was originally associated: *Begomovirus brassicae*. |
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| **Permission for use of names derived from a living person:**  |
| **Taxon name** | **Full name of person from whom the name is derived** | **Attached**  |
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| **Abstract of Taxonomy Proposal:**  |
| *Taxonomic rank affected*:Genus *Deltasatellite* in the family *Tolecusatellitidae* *Description of current taxonomy*:*Tolecusatellitidae /* *Deltasatellite*The genus *Deltasatellite* consists of 12 species.*Proposed* *taxonomic change:*We proposed to create one new species in the genus *Deltasatellite*: “*Deltasatellite brassicae*”.*Justification*:Similar to members of the genus *Deltasatellite*, members of the proposed new species “*Deltasatellite brassicae*” have all conserved features of deltasatellites, including an A-rich region, a stem-loop with the TAATATTAC nonanucleotide motif, and a region with identity to betasatellites. The isolates share 100% nucleotide identity among themselves, and ≤73.9% identity with known deltasatellites, below the 91% species demarcation threshold. Phylogenetic analysis shows the group forms a distinct clade within New World deltasatellites. We conclude that “*Deltasatellite brassicae*”represents a new species in the genus *Deltasatellite*. |

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| *Taxonomic rank affected:* Genus *Deltasatellite* in the family *Tolecusatellitidae* *Description of current taxonomy:* *Tolecusatellitidae /Deltasatellite*The genus *Deltasatellite* consists of 12 species.Twelve species are currently assigned to the *Deltasatellite* genus in the *Tolecusatellitidae* family:*Deltasatellite codiaeumiflavi, Deltasatellite desmodii, Deltasatellite ipomoeaprimi, Deltasatellite ipomoeasecundi, Deltasatellite ipomoeatertii, Deltasatellite malvastri, Deltasatellite sidaflavusprimi, Deltasatellite sidaflavussecundi, Deltasatellite sidaflavustertii, Deltasatellite solani, Deltasatellite solaniflavusprimi, Deltasatellite solaniflavussecundi.**Proposed taxonomic change:* We proposed to create one new species in the genus *Deltasatellite*: “*Deltasatellite brassicae*”, which was isolated associated to the begomovirus cabbage leaf curl virus (*Begomovirus brassicae*) from black gram (*Vigna mungo*), cowpea (*Vigna unguiculata*), *Desmodium scorpiurus*, and *Rhynchosia minima* collected from Venezuela [1]. The satellite was termed cabbage leaf curl deltasatellite (CabLCD).*Demarcation criteria:*A 91% pairwise identity species demarcation threshold has been proposed and adopted for the genus *Deltasatellite* [2].*Justification:* The sequences of CabLCD summarized in Table 1 can be classified into one new species sharing ≤73.9% genome-wide pairwise identity with all classified deltasatellites (Figure 1). Furthermore, this is supported by phylogenetic analysis (Figure 2). |

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| **References:**  |
| [1] Fiallo-Olivé, E., Bastidas, L., Chirinos, D.T., Navas-Castillo, J., 2021. Insights into emerging begomovirus-deltasatellite complex diversity: the first deltasatellite infecting legumes. Biology 10, 1125.[2] Briddon, R.W., Navas-Castillo, J., Fiallo-Olivé, E., 2016. ICTV taxonomic Proposal 2016.021a-kP.A.v2.Tolecusatellitidae. Create the *Tolecusatellitidae*, a new family of single-stranded DNA satellites with two genera. Available online at: http://www.ictv.global/proposals-16/2016.021a-kP.A.v2.Tolecusatellitidae.pdf (accessed June 1, 2025).[3] Muhire, B.M., Varsani, A., Martin, D.P., 2014. SDT: a virus classification tool based on pairwise sequence alignment and identity calculation. *PloS one* *9*, e108277. |

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| **Accompanying files:**  |
| **Filename** | **Description of contents** |
| 2025.015P.A.v1.Deltasatellite\_1nsp | spreadsheet |
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| **Tables, Figures:**  |

**Table 1:** Summary of the new proposed species in the *Deltasatellite* genus of the *Tolecusatellitidae* family. The exemplar isolate is highlighted in red.

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| **Genus** | **Species** | **Accession #** | **Deltasatellite name** | **Acronym** | **Isolate** | **Country** | **Host** | **Associated begomovirus species** |
| *Deltasatellite* | *“Deltasatellite brassicae”* | OK073969 | Cabbage leaf curl deltasatellite | CabLCD | VE-Cp\_V1-17 | Venezuela | Cowpea | *Begomovirus brassicae* |
| *Deltasatellite* | *Deltasatellite brassicae* | OK073970 | Cabbage leaf curl deltasatellite | CabLCD | VE-Cp\_V2-17 | Venezuela | Cowpea | *Begomovirus brassicae* |
| *Deltasatellite* | *Deltasatellite brassicae* | OK073971 | Cabbage leaf curl deltasatellite | CabLCD | VE-Des\_V8-17 | Venezuela | *Desmodium scorpiurus* | *Begomovirus brassicae* |
| *Deltasatellite* | *Deltasatellite brassicae* | OK073972 | Cabbage leaf curl deltasatellite | CabLCD | VE-Rh\_V9-17 | Venezuela | *Rhynchosia minima* | *Begomovirus brassicae* |
| *Deltasatellite* | *Deltasatellite brassicae* | OK073973 | Cabbage leaf curl deltasatellite | CabLCD | VE-Rh\_V10-17 | Venezuela | *Rhynchosia minima* | *Begomovirus brassicae* |
| *Deltasatellite* | *Deltasatellite brassicae* | OK073974 | Cabbage leaf curl deltasatellite | CabLCD | VE-Bg\_V12-17 | Venezuela | Black gram | *Begomovirus brassicae* |
| *Deltasatellite* | *Deltasatellite brassicae* | OK073975 | Cabbage leaf curl deltasatellite | CabLCD | VE-Bg\_V13-17 | Venezuela | Black gram | *Begomovirus brassicae* |



**Figure 1:** Pairwise identity matrix inferred using SDT [3]. A representative sequence of the proposed new species is highlighted in red.



**Figure 2:** Maximum likelihood phylogenetic tree of representative genomes from each deltasatellite species. Bootstrap values are shown at the nodes. A representative sequence of the proposed new species is highlighted in red.