This form should be used for all taxonomic proposals. Please complete all those modules that are applicable (and then delete the unwanted sections). For guidance, see the notes written in blue and the separate document “Help with completing a taxonomic proposal”

Please try to keep related proposals within a single document; you can copy the modules to create more than one genus within a new family, for example.

**MODULE 1: TITLE, AUTHORS, etc**

<table>
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<th>2015.006aB</th>
<th>(to be completed by ICTV officers)</th>
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<tbody>
<tr>
<td><strong>Short title:</strong></td>
<td>Renaming of all phage genera (e.g. 6 new species in the genus Zetavirus)</td>
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<td><strong>Modules attached</strong></td>
<td>1 ☒</td>
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<tr>
<td>(modules 1 and 10 are required)</td>
<td>6 ☒</td>
<td>7 ☒</td>
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**Author(s):**

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**List the ICTV study group(s) that have seen this proposal:**

| A list of study groups and contacts is provided at [http://www.ictvonline.org/subcommittees.asp](http://www.ictvonline.org/subcommittees.asp). If in doubt, contact the appropriate subcommittee chair (fungal, invertebrate, plant, prokaryote or vertebrate viruses) | Bacterial and Archaeal Viruses Subcommittee |

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The Bacterial and Archaeal Virus Subcommittee of ICTV has voted overwhelmingly in favour of eliminating “like” and “Phi” from phage genus names. The latter largely takes effect with the establishment of new genera.
**MODULE 8: RENAME**

Use this module to change the name of one or more existing taxa (but note that stability of nomenclature is encouraged wherever possible). Insert extra lines in the table if needed.

Renaming one or more taxa

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Reasons to justify the renaming:

Explain why the taxon (or taxa) should be renamed

The Bacterial and Archaeal Virus Subcommittee has unanimously voted to no longer use “like” and “phi” in the names of bacteriophage genera. The latter largely takes effect with the establishment of new genera.

N.B. 1  Genus name non-standard
N.B. 2  Name difficult to pronounce.
N.B. 3. Renamed so as to reduce confusion with Hpi1virus; and Pbi1virus, respectively
N.B. 4  Phage I3 has not be completely sequenced, therefore named after first completely sequenced phage of this type
N.B. 5 This is an archaeal virus; renamed by the Study Group Chair, David Prangishvili