



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**

Code assigned:	2016.076aB	(to be completed by ICTV officers)			
Short title: To rename thirteen (13) existing species and one (1) genus. (e.g. 6 new species in the genus <i>Zetavirus</i>)					
Modules attached (modules 1 and 11 are required)	1 <input checked="" type="checkbox"/> 6 <input type="checkbox"/>	2 <input type="checkbox"/> 7 <input type="checkbox"/>	3 <input type="checkbox"/> 8 <input type="checkbox"/>	4 <input type="checkbox"/> 9 <input checked="" type="checkbox"/>	5 <input type="checkbox"/> 10 <input type="checkbox"/>

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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> . If in doubt, contact the appropriate subcommittee chair (fungal, invertebrate, plant, prokaryote or vertebrate viruses)

ICTV Bacterial and Archaeal Viruses Subcommittee

ICTV Study Group comments (if any) and response of the proposer:

Date first submitted to ICTV:

July 2016

Date of this revision (if different to above):

21 Sept 2016

ICTV-EC comments and response of the proposer:

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MODULE 9: **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

Code	2016.076aB	(assigned by ICTV officers)
To rename the following taxon (or taxa):		
Current name	Proposed name	
<i>Haloarcula virus HHPV-1</i>	<i>Haloarcula virus HHPV1</i>	
<i>Haloarcula virus HHPV-2</i>	<i>Haloarcula virus HHPV2</i>	
<i>Halorubrum virus HRPV-1</i>	<i>Halorubrum virus HRPV1</i>	
<i>Halorubrum virus HRPV-2</i>	<i>Halorubrum virus HRPV2</i>	
<i>Halorubrum virus HRPV-6</i>	<i>Halorubrum virus HRPV6</i>	
<i>Halogeometricum virus HGPV-1</i>	<i>Halogeometricum virus HGPV1</i>	
<i>Halorubrum virus HRPV-3</i>	<i>Halorubrum virus HRPV3</i>	
<i>Mannheimia virus MhaA1- PHL101</i>	<i>Mannheimia virus PHL101</i>	
<i>Enterobacter virus CC31</i>	<i>Escherichia virus CC31</i>	
<i>Lactobacillus phage Lb338-1</i>	<i>Lactobacillus virus Lb338-1</i>	
<i>Mycobacterium virus Fionn</i>	<i>Mycobacterium virus Fionnbarth</i>	
<i>Salmonella phage L13</i>	<i>Salmonella virus L13</i>	
<i>Salmonella phage LSPA1</i>	<i>Salmonella virus LSPA1</i>	
<i>Bcep22likevirus</i>	<i>Bcep22virus</i>	

Reasons to justify the renaming:

Explain why the taxon (or taxa) should be renamed

In TaxoProp 2015.025aB “To rename all (522) existing bacterial virus and 2 archaeal virus species” we stated “(f) remove hyphens from the virus species names in accordance with the ICVCN rule 3.13. N.B. Hyphens between two numbers or two letters were retained to avoid confusion and loss of encoded information.” Therefore, the hyphen was removed for the first seven taxa. In the case of the latter four phages the word “phage” was replaced with “virus”; and, lastly, the actual host of CC31 is Escherichia not Enterobacter. In the case of the Mycobacterium virus the correct name for the species is Fionnbarth. “Like” has been abolished from all other genera names.