



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:	2014.016aV	(to be completed by ICTV officers)
Short title: Rename 12 picornavirus species (e.g. 6 new species in the genus <i>Zetavirus</i>)		
Modules attached (modules 1 and 9 are required)	1 <input checked="" type="checkbox"/> 2 <input type="checkbox"/> 3 <input type="checkbox"/> 4 <input type="checkbox"/> 5 <input type="checkbox"/> 6 <input type="checkbox"/> 7 <input type="checkbox"/> 8 <input checked="" type="checkbox"/> 9 <input checked="" type="checkbox"/>	

Author(s) with e-mail address(es) of the proposer:

Nick Knowles (nick.knowles@pirbright.ac.uk) on behalf of the *Picornaviridae* Study Group

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)

Picornaviridae SG

ICTV-EC or Study Group comments and response of the proposer:

Date first submitted to ICTV:

07/07/2014

Date of this revision (if different to above):

MODULE 8: **NON-STANDARD**

Template for any proposal not covered by modules 2-7. This includes proposals to change the name of existing taxa (but note that stability of nomenclature is encouraged wherever possible).

non-standard proposal

Code	2014.016aV	(assigned by ICTV officers)
Title of proposal: Renaming of 12 picornavirus species		

Text of proposal:

In order to move towards a more consistent genus/species nomenclature (and in some cases to remove host species references) it is proposed to rename the species shown in Table 1. The virus common names, e.g. encephalomyocarditis virus would remain unchanged, thus helping to distinguish them from the species names. Possible renaming members of the *Aphthovirus* genus will be addressed in the future.

Table 1. Proposed new picornavirus species names.

Genus	Current species name	Proposed new species name	Virus common name*
<i>Avihepatovirus</i>	<i>Duck hepatitis A virus</i>	<i>Avihepatovirus A</i>	duck hepatitis A virus 1-3
<i>Cardiovirus</i>	<i>Encephalomyocarditis virus</i>	<i>Cardiovirus A</i>	encephalomyocarditis virus 1-2
<i>Cardiovirus</i>	<i>Theilovirus</i>	<i>Cardiovirus B</i>	Theiler's murine encephalomyelitis virus, Vilyuisk human encephalomyelitis virus, thera virus, Saffold virus 1-11
<i>Erbovirus</i>	<i>Equine rhinitis B virus</i>	<i>Erbovirus A</i>	equine rhinitis B virus 1-3
<i>Hepatovirus</i>	<i>Hepatitis A virus</i>	<i>Hepatovirus A</i>	hepatitis A virus 1
<i>Parechovirus</i>	<i>Human parechovirus</i>	<i>Parechovirus A</i>	human parechovirus 1-16
<i>Parechovirus</i>	<i>Ljungan virus</i>	<i>Parechovirus B</i>	Ljungan virus 1-4
<i>Sapelovirus</i>	<i>Porcine sapelovirus</i>	<i>Sapelovirus A</i>	porcine sapelovirus 1
<i>Sapelovirus</i>	<i>Simian sapelovirus</i>	<i>Sapelovirus B</i>	simian sapelovirus 1-3
<i>Senecavirus</i>	<i>Seneca Valley virus</i>	<i>Senecavirus A</i>	Seneca Valley virus 1
<i>Teschovirus</i>	<i>Porcine teschovirus</i>	<i>Teschovirus A</i>	porcine teschovirus 1-13
<i>Tremovirus</i>	<i>Avian encephalomyelitis virus</i>	<i>Tremovirus A</i>	avian encephalomyelitis virus 1

*, the virus common name would remain unchanged thus differentiating it from the species name.

MODULE 9: **APPENDIX**: supporting material

additional material in support of this proposal

References:

None.

Annex:

Include as much information as necessary to support the proposal, including diagrams comparing the old and new taxonomic orders. The use of Figures and Tables is strongly recommended but direct pasting of content from publications will require permission from the copyright holder together with appropriate acknowledgement as this proposal will be placed on a public web site. For phylogenetic analysis, try to provide a tree where branch length is related to genetic distance.
