Ratification vote on taxonomic proposals to the International Committee on Taxonomy of Viruses (2015)

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Abstract Changes to virus taxonomy approved and ratified by the International Committee on Taxonomy of Viruses in February 2015 are listed.

Changes to virus taxonomy (the Universal Scheme of Virus Classification of the International Committee on Taxonomy of Viruses [ICTV]) now take place annually and are the result of a multi-stage process. In accordance with the ICTV Statutes, proposals submitted to the Executive Committee undergo a review process that involves input from the ICTV Study Groups (SGs) and Subcommittees (SCs), other interested virologists, and the ICTV Executive Committee. Reviewed proposals are then presented for ratification to the full ICTV membership by publication on an ICTV web site followed by an electronic vote. The latest list of approved proposals was prepared and presented on the ICTV online web pages by November 2014. This list of proposals was then sent via email on 20 January 2015 to the 146 members of ICTV, namely the ICTV Executive Committee Members, Life Members, ICTV Subcommittee Members and ICTV National Representatives. Members were then requested to vote on whether or not to ratify the taxonomic proposals (voting closed on 20 February 2015).

This report lists the taxonomic proposals that were ratified by ICTV members. Note that by ICTV convention, all approved taxon names are italicized. The

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proposals are listed by Subcommittee, and then the file-name as posted on-line. For the first time, each proposal is formally cited so that a list of the authors appears in the References. This provides a means of acknowledging their efforts and also links to the specific proposal on the ICTV web site.

Invertebrate virus subcommittee

2013.002aI.A.v3.Betaentomopoxvirus_3sp [1]

2013.002aI   create 3 species, *Adoxophyes honmai entomopoxvirus*, *Choristoneura rosaceana entomopoxvirus* and *Mythimna separata entomopoxvirus* in the genus *Betaentomopoxvirus*, subfamily *Entomopoxvirinae*, family *Poxviridae*

2014.001aI.A.v1. Entomopoxvirus-spcorr [2]

2014.001aI   Correct typographical and other errors in species names in the subfamily *Entomopoxvirinae*:

Old name	New name
(a) in genus *Alphaentomopoxvirus*:	
Figulus subleavis entomopoxvirus	*Figulus sublaevis entomopoxvirus*
Demodema boranensis entomopoxvirus	*Demodema bonariensis entomopoxvirus*
(b) in genus *Betaentomopoxvirus*:	
Acrobasis zelleri entomopoxvirus 'L'	*Acrobasis zelleri entomopoxvirus*
Amsacta moorei entomopoxvirus 'L'	*Amsacta moorei entomopoxvirus*
Arphia conspersa entomopoxvirus 'O'	*Arphia conspersa entomopoxvirus*
Choristoneura biennis entomopoxvirus 'L'	*Choristoneura biennis entomopoxvirus*
Choristoneura conflicta entomopoxvirus 'L'	*Choristoneura conflicta entomopoxvirus*
Choristoneura diversuma entomopoxvirus 'L'	*Choristoneura diversuma entomopoxvirus*
Choristoneura fumiferana entomopoxvirus 'L'	*Choristoneura fumiferana entomopoxvirus*
Chorizagrotis auxiliars entomopoxvirus 'L'	*Chorizagrotis auxiliaris entomopoxvirus*
Heliothis armigera entomopoxvirus 'L'	*Heliothis armigera entomopoxvirus*
Locusta migratoria entomopoxvirus 'O'	*Locusta migratoria entomopoxvirus*
Oedaleus senigalensis entomopoxvirus 'O'	*Oedaleus senegalensis entomopoxvirus*
Operophtera brumata entomopoxvirus 'L'	*Operophtera brumata entomopoxvirus*
Schistocera gregaria entomopoxvirus 'O'	*Schistocerca gregaria entomopoxvirus*
(c) in genus *Gammaentomopoxvirus*:	
Goeldichironomus haloprasimus entomopoxvirus	*Goeldichironomus holoprasinus entomopoxvirus*

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2014.002aI.A.v1. Alphabaculovirus-sprem [3]

2014.002aI		remove species *Helicoverpa zea single nucleopolyhedrovirus* from the genus *Alphabaculovirus*

Plant virus subcommittee

2013.010a,bP.A.v2.Umbravirus_move [4]

2013.010aP		remove genus *Umbravirus* from its present (unassigned) position

2013.010bP		reassign genus *Umbravirus* to the family *Tombusviridae*

2014.001aP.A.v1.Tobamovirus_2sp [5]

2014.001aP		create 2 species, *Tomato mottle mosaic virus* and *Yellow tailflower mild mottle virus*, in the genus *Tobamovirus*, family *Virgaviridae*

2014.002aP.A.v1.Pospiviroid_sprem [6]

2014.002aP		remove (abolish) the species *Mexican papita viroid* (currently in the genus *Pospiviroid*, family *Pospiviroidae*)

2014.003aP.A.v2.Hostuviroid_sp [7]

2014.003aP		create species *Dahlia latent viroid* in the genus *Hostuviroid*

2014.004aP.A.v2.Torradovirus_sp [8]

2014.004aP		create species *Lettuce necrotic leaf curl virus* in the genus *Torradovirus*, family *Secoviridae*, order *Picornavirales*

2014.005aP.A.v1.Tritimovirus_sp [9]

2014.005aP		create the species *Tall oatgrass mosaic virus* in the genus *Tritimovirus*, family *Potyviridae*

2014.006aP.A.v2.Tombusviridae_4sp [10]

2014.006aP		create 4 species (*Elderberry latent virus*, *Pelargonium chlorotic ring pattern virus*, *Pelargonium ringspot virus* and *Rosa rugosa leaf distortion virus*) to be unassigned in the family *Tombusviridae*

2014.007aP.A.v1.Panicovirus_sp [11]

2014.007aP		create species *Thin paspalum asymptomatic virus* in genus *Panicovirus* in the family *Tombusviridae*

2014.008aP.A.v3.Tombusviridae_sp [12]

2014.006aP		create species *Trailing lespedeza virus 1* to be unassigned in the family *Tombusviridae*

2014.009aP.A.v4.Potyvirus_12sp [13]

2014.009aP		create 12 species (*Bidens mosaic virus*, *Blue squill virus A*, *Brugmansia mosaic virus*, *Calla lily latent virus*, *Habenaria mosaic virus*, *Keunjorong mosaic virus*, *Lupinus mosaic virus*, *Panax virus Y*, *Tomato necrotic stunt virus*, *Cyrtanthus elatus virus A*, *Verbena virus Y* and *Zucchini tigre mosaic virus*) in the genus *Potyvirus*, family *Potyviridae*

2014.010a-dP.A.v2.Rosadnavirus [14]

2014.010aP		create species *Rose yellow vein virus* in genus *Rosadnavirus* (new) in the family *Caulimoviridae*

2014.010bP		create a new genus

2014.010cP		name the genus *Rosadnavirus*

2014.010dP		designate *Rose yellow vein virus* as type species of the new genus

2014.011aP.A.v1.Caulimovirus_sp [15]

2014.011aP		create species *Soybean Putnam virus* in genus *Caulimovirus*, family *Caulimoviridae*

2014.012aP.A.v1.Badnavirus_7sp [16]

2014.012aP		create 7 species (*Banana streak IM virus*, *Banana streak UA virus*, *Banana streak UI virus*, *Banana streak UL virus*, *Banana streak UM virus*, *Fig badnavirus 1* and *Pagoda yellow mosaic associated virus*) in the genus *Badnavirus*, family *Caulimoviridae*

2014.013aP.A.v2.Emaravirus_sp [17]

2014.013aP		create species *Pigeonpea sterility mosaic virus* in the genus *Emaravirus*

Prokaryote virus subcommittee

2012.002a-dB.A.v3.Corndoglikevirus [18]

2012.002aB		create 2 species (*Mycobacterium phage corndog* and *Mycobacterium phage firecracker*) in the genus *Corndoglikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2012.002bB		create a new genus in the family *Siphoviridae*

2012.002cB		name the new genus *Corndoglikevirus*

2012.002dB		designate *Mycobacterium phage corndog* as the type species of the new genus

2012.003a-dB.A.v3.D3112likevirus [19]

2012.003aB

create 8 species (*Pseudomonas phage D3112*, *Pseudomonas phage Dms3*, *Pseudomonas phage Fha0480*, *Pseudomonas phage Lpb1*, *Pseudomonas phage Mp22*, *Pseudomonas phage Mp29*, *Pseudomonas phage Mp38* and *Pseudomonas phage Pa1phi*) in the genus *D3112likevirus* (new), family *Siphoviridae*, order *Caudovirales*

2012.003bB   create a new genus in the family *Siphoviridae*

2012.003cB   name the new genus *D3112likevirus*

2012.003dB   designate *Pseudomonas phage D3112* as the type species of the new genus

2013.001a-oB.A.v5.Sphaerolipoviridae [20, 21]

2013.001aB   create 3 species (*Haloarcula hispanica virus SH1*, *Haloarcula hispanica virus PH1* and *Haloarcula hispanica icosahedral virus 2*) in the genus *Alphasphaerolipovirus* (new), family *Sphaerolipoviridae* (new)

2013.001bB   create a new genus in the family *Sphaerolipoviridae* (new)

2013.001cB   name the new genus *Alphasphaerolipovirus*

2013.001dB   designate *Haloarcula hispanica virus SH1* as the type species of the new genus

2013.001eB   create species *Natrinema virus SNJ1* in the genus *Betasphaerolipovirus* (new), family *Sphaerolipoviridae* (new)

2013.001fB   create a new genus in the family *Sphaerolipoviridae* (new)

2013.001gB   name the new genus *Betasphaerolipovirus*

2013.001hB   designate *Natrinema virus SNJ1* as the type species of the new genus

2013.001iB   create 2 species, *Thermus thermophilus phage P23-77* and *Thermus thermophilus phage IN93*, in the genus *Gammasphaerolipovirus* (new), family *Sphaerolipoviridae* (new)

2013.001jB   create a new genus in the family *Sphaerolipoviridae* (new)

2013.001kB   name the new genus *Gammasphaerolipovirus*

2013.001lB   designate *Thermus thermophilus phage P23-77* as the type species of the new genus

2013.001mB   create a new unassigned family

2013.001nB   name the new family *Sphaerolipoviridae*

2013.001oB   assign 3 genera, *Alphasphaerolipovirus*, *Betasphaerolipovirus* and *Gammasphaerolipovirus*, to the new family

2013.004a-kB.A.v4.Eucampyvirinae [22, 23]

2013.004aB   create 4 species (*Campylobacter phage CP220*, *Campylobacter phage CPt10*, *Campylobacter phage IBB35* and *Campylobacter phage CP21*) in the genus *Cp220likevirus* (new), subfamily *Eucampyvirinae* (new), family *Myoviridae*, order *Caudovirales*

2013.004bB   create 3 species (*Campylobacter phage CP81*, *Campylobacter phage CPX* and *Campylobacter phage NCTC12673*) in the genus *Cp8unalikevirus* (new), subfamily *Eucampyvirinae* (new), family *Myoviridae*, order *Caudovirales*

2013.004cB   create a new genus in the subfamily *Eucampyvirinae* (new), family *Myoviridae*

2013.004dB   name the new genus *Cp220likevirus*

2013.004eB   designate *Campylobacter phage CP220* as the type species of the new genus

2013.004fB   create a new genus in the subfamily *Eucampyvirinae* (new), family *Myoviridae*

2013.004gB   name the new genus *Cp8unalikevirus*

2013.004hB   designate *Campylobacter phage CP81* as the type species of the new genus

2013.004iB   create a new subfamily in the family *Myoviridae*

2013.004jB   name the new subfamily *Eucampyvirinae*

2013.004kB   assign 2 genera, *Cp220likevirus* and *Cp8unalikevirus*, to the new subfamily

2013.005a-dB.A.v3.Phifllikevirus [24]

2013.005aB   create 3 species (*Enterococcus phage phifl1*, *Enterococcus phage phifl2 and Enterococcus phage phifl3*) in the genus *Phifllikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.005bB   create a new genus in the family *Siphoviridae*

2013.005cB   name the new genus *Phifllikevirus*

2013.005dB   designate *Enterococcus phage phifl1* as the type species of the new genus

2013.006a-dB.A.v3.Iebhlikevirus [25]

2013.006aB   create 2 species, *Bacillus phage IEBH* and *Bacillus phage 250*, in the genus *Iebhlikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.006bB   create a new genus in the family *Siphoviridae*

2013.006cB   name the new genus *Iebhlikevirus*

2013.006dB   designate *Bacillus phage IEBH* as the type species of the new genus

2013.007a-dB.A.v3.Halolikevirus [26]

2013.007aB   create 2 species, *Mycobacterium phage halo* and *Mycobacterium phage liefie*, in the genus *Halolikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.007bB   create a new genus in the family *Siphoviridae*

2013.007cB   name the new genus *Halolikevirus*

2013.007dB   designate *Mycobacterium phage halo* as the type species of the new genus

2013.008a-dB.A.v3.Che8likevirus [27]

2013.008aB   create the following 28 species in the genus *Che8likevirus* (new), family *Siphoviridae*, order *Caudovirales*: *Mycobacterium phage Che8*, *Mycobacterium phage PMC*, *Mycobacterium phage mutaforma13*, *Mycobacterium phage dotproduct*, *Mycobacterium phage gumbie*, *Mycobacterium phage Llij*, *Mycobacterium phage spartacus*, *Mycobacterium phage Pacc40*, *Mycobacterium phage ibhubesi*, *Mycobacterium phage tweety*, *Mycobacterium phage taj*, *Mycobacterium phage wee*, *Mycobacterium phage dorothy*, *Mycobacterium phage deadp*, *Mycobacterium phage SG4*, *Mycobacterium phage boomer*, *Mycobacterium phage dlane*, *Mycobacterium phage ardmore*, *Mycobacterium phage fruitloop*, *Mycobacterium phage shilan*, *Mycobacterium phage drago*, *Mycobacterium phage ramsey*, *Mycobacterium phage rockyhorror*, *Mycobacterium phage mozy*, *Mycobacterium phage shauna1*, *Mycobacterium phage Che9d*, *Mycobacterium phage avani* and *Mycobacterium phage yoshi*

2013.008bB   create a new genus in the family *Siphoviridae*

2013.008cB   name the new genus *Che8likevirus*

2013.008dB   designate *Mycobacterium phage Che8* as the type species of the new genus

2013.009a-dB.A.v3.Bignuzlikevirus [28]

2013.009aB   create 2 species, *Mycobacterium phage bignuz* and *Mycobacterium phage jebeks*, in the genus *Bignuzlikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.009bB   create a new genus in the family *Siphoviridae*

2013.009cB   name the new genus *Bignuzlikevirus*

2013.009dB   designate *Mycobacterium phage bignuz* as the type species of the new genus

2013.010a-dB.A.v3.Phijlunalikevirus [29]

2013.010aB   create 3 species (*Lactobacillus phage phijl1*, *Pediococcus phage clp1* and *Lactobacillus phage ATCC8014*) in the genus *Phijlunalikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.010bB   create a new genus in the family *Siphoviridae*

2013.010cB   name the new genus *Phijlunalikevirus*

2013.010dB   designate *Lactobacillus phage phijl1* as the type species of the new genus

2013.011a-dB.A.v3.Barnyardlikevirus [30]

2013.011aB   create 4 species (*Mycobacterium phage barnyard*, *Mycobacterium phage predator*, *Mycobacterium phage konstantine* and *Mycobacterium phage patience*) in the genus *Barnyardlikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.011bB   create a new genus in the family *Siphoviridae*

2013.011cB   name the new genus *Barnyardlikevirus*

2013.011dB   designate *Mycobacterium phage barnyard* as the type species of the new genus

2013.012a-dB.A.v3.Charlielikevirus [31]

2013.012aB   create 2 species, *Mycobacterium phage charlie* and *Mycobacterium phage redi*, in the genus *Charlielikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.012bB   create a new genus in the family *Siphoviridae*

2013.012cB   name the new genus *Charlielikevirus*

2013.012dB   designate *Mycobacterium phage charlie* as the type species of the new genus

2013.013a-dB.A.v3.Che9clikevirus [32]

2013.013aB   create 3 species (*Mycobacterium phage Che9c*, *Mycobacterium phage brujita* and *Mycobacterium phage babsiella*) in the genus *Che9clikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.013bB   create a new genus in the family *Siphoviridae*

2013.013cB   name the new genus *Che9clikevirus*

2013.013dB   designate *Mycobacterium phage Che9c* as the type species of the new genus

2013.014a-dB.A.v3.Cjwunalikevirus [33]

2013.014aB   create 9 species (*Mycobacterium phage CJW1*, *Mycobacterium phage pumpkin*, *Mycobacterium phage 244*, *Mycobacterium phage porky*, *Mycobacterium phage Bask21*, *Mycobacterium phage sirduracell*, *Mycobacterium phage eureka*, *Mycobacterium phage kostya* and *Mycobacterium phage toto*) in the genus *Cjwunalikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.014bB   create a new genus in the family *Siphoviridae*

2013.014cB   name the new genus *Cjwunalikevirus*

2013.014dB   designate *Mycobacterium phage CJW1* as the type species of the new genus

2013.015a-dB.A.v3.D3likevirus [34]

2013.015aB   create 2 species, *Pseudomonas phage D3* and *Pseudomonas phage PMG1*, in the genus *D3likevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.015bB	create a new genus in the family *Siphoviridae*
2013.015cB	name the new genus *D3likevirus*
2013.015dB	designate *Pseudomonas phage D3* as the type species of the new genus

2013.016aB.A.v3.L5likevirus_62sp [35]

2013.016aB   create the following 62 species in the genus *L5likevirus*, family *Siphoviridae*, order Caudovirales: *Mycobacterium phage L5*, *Mycobacterium phage SWU1*, *Mycobacterium phage Che12*, *Mycobacterium phage D29*, *Mycobacterium phage trixie*, *Mycobacterium phage turbido*, *Mycobacterium phage Ta17a*, *Mycobacterium phage redrock*, *Mycobacterium phage pukovnik*, *Mycobacterium phage ericb*, *Mycobacterium phage hammer*, *Mycobacterium phage gladiator*, *Mycobacterium phage jeffabunny*, *Mycobacterium phage Bxb1*, *Mycobacterium phage billknuckles*, *Mycobacterium phage switzer*, *Mycobacterium phage kssjeb*, *Mycobacterium phage U2*, *Mycobacterium phage mrgordo*, *Mycobacterium phage museum*, *Mycobacterium phage lesedi*, *Mycobacterium phage solon*, *Mycobacterium phage bethlehem*, *Mycobacterium phage KBG*, *Mycobacterium phage kugel*, *Mycobacterium phage bruns*, *Mycobacterium phage violet*, *Mycobacterium phage euphoria*, *Mycobacterium phage BPBiebs31*, *Mycobacterium phage nepal*, *Mycobacterium phage doom*, *Mycobacterium phage marcell*, *Mycobacterium phage lockley*, *Mycobacterium phage skipole*, *Mycobacterium phage JC27*, *Mycobacterium phage Jasper*, *Mycobacterium phage Ridgecb*, *Mycobacterium phage Perseus*, *Mycobacterium phage Bxz2*, *Mycobacterium phage JHC117*, *Mycobacterium phage microwolf*, *Mycobacterium phage wonder*, *Mycobacterium phage heldan*, *Mycobacterium phage rockstar*, *Mycobacterium phage peaches*, *Mycobacterium phage arturo*, *Mycobacterium phage LHTSCC*, *Mycobacterium phage backyardigan*, *Mycobacterium phage benedict*, *Mycobacterium phage george*, *Mycobacterium phage cuco*, *Mycobacterium phage tiger*, *Mycobacterium phage timshel*, *Mycobacterium phage saintus*, *Mycobacterium phage astro*, *Mycobacterium phage packman*, *Mycobacterium phage alma*, *Mycobacterium phage rebeuca*, *Mycobacterium phage goose*, *Mycobacterium phage twister*, *Rhodococcus phage RGL3* and *Rhodococcus phage RER2*

2013.017a-dB.A.v3.Bronlikevirus [36]

2013.017aB	create 4 species (*Mycobacterium phage bron*, *Mycobacterium phage joedirt*, *Mycobacterium phage rumpelstiltskin* and *Mycobacterium phage Faith1*) in the genus *Bronlikevirus* (new), family *Siphoviridae*, order *Caudovirales*
2013.017bB	create a new genus in the family *Siphoviridae*
2013.017cB	name the new genus *Bronlikevirus*
2013.017dB	designate *Mycobacterium phage bron* as the type species of the new genus

2013.018a-dB.A.v3.Omegalikevirus [37]

2013.018aB	create 6 species (*Mycobacterium phage omega*, *Mycobacterium phage littlee*, *Mycobacterium phage courthouse*, *Mycobacterium phage optimus*, *Mycobacterium phage baka and Mycobacterium phage thibault*) in the genus *Omegalikevirus* (new), family *Siphoviridae*, order *Caudovirales*
2013.018bB	create a new genus in the family *Siphoviridae*
2013.018cB	name the new genus *Omegalikevirus*
2013.018dB	designate *Mycobacterium phage omega* as the type species of the new genus

2013.019a-dBA.v4.Pbiunalikevirus [38]

2013.019aB	create species *Mycobacterium phage PBI1* in the genus *Pbiunalikevirus* (new), family *Siphoviridae*, order *Caudovirales*
2013.019bB	create a new genus in the family *Siphoviridae*
2013.019cB	name the new genus *Pbiunalikevirus*
2013.019dB	designate *Mycobacterium phage PBI1* as the type species of the new genus

2013.020a-dB.A.v4.Pgonelikevirus [39]

2013.020aB	create 12 species (*Mycobacterium phage Pg1*, *Mycobacterium phage oline*, *Mycobacterium phage rosebush*, *Mycobacterium phage pipefish*, *Mycobacterium phage gadjet*, *Mycobacterium phage athena*, *Mycobacterium phage cooper*, *Mycobacterium phage chrisnmich*, *Mycobacterium phage stinger*, *Mycobacterium phage nigel*, *Mycobacterium phage zemanar* and *Mycobacterium phage acadian*) in the genus *Pgonelikevirus* (new), family *Siphoviridae*, order *Caudovirales*
2013.020bB	create a new genus in the family *Siphoviridae*
2013.020cB	name the new genus *Pgonelikevirus*
2013.020dB	designate *Mycobacteriophage Pg1* as the type species of the new genus

2013.021a-dB.A.v3.Reylikevirus [40]

2013.021aB	create 2 species, *Mycobacterium phage rey* and *Mycobacterium phage bongo*, in the genus *Reylikevirus* (new), family *Siphoviridae*, order *Caudovirales*
2013.021bB	create a new genus in the family *Siphoviridae*
2013.021cB	name the new genus *Reylikevirus*
2013.021dB	designate *Mycobacterium phage rey* as the type species of the new genus

2013.022a-dB.A.v3.Tm4likevirus [41]

2013.022aB	create 9 species (*Mycobacterium phage TM4*, *Mycobacterium phage angelica*, *Mycobacterium phage jaws*, *Mycobacterium phage crimd*, *Mycobacterium phage anaya*, *Mycobacterium phage pixie*, *Mycobacterium phage macncheese*, *Mycobacterium phage larva* and *Mycobacterium phage fionn*) in the genus *Tm4likevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.022bB create a new genus in the family *Siphoviridae*

2013.022cB name the new genus *Tm4likevirus*

2013.022dB designate *Mycobacterium phage TM4* as the type species of the new genus

2013.023a-dB.A.v3.C5likevirus [42]

2013.023aB create 2 species, *Lactobacillus phage c5* and *Lactobacillus phage LLKu*, in the genus *C5likevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.023bB create a new genus in the family *Siphoviridae*

2013.023cB name the new genus *C5likevirus*

2013.023dB designate *Lactobacillus phage c5* as the type species of the new genus

2013.024a-dB.A.v3.P23likevirus [43]

2013.024aB create 2 species, *Thermus phage P2345* and *Thermus phage P7426*, in the genus *P23likevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.024bB create a new genus in the family *Siphoviridae*

2013.024cB name the new genus *P23likevirus*

2013.024dB designate *Thermus phage P2345* as the type species of the new genus

2013.025a-dB.A.v3.Phicbklikevirus [44]

2013.025aB create 5 species (*Caulobacter phage phicbk*, *Caulobacter phage magneto*, *Caulobacter phage swift*, *Caulobacter phage karma* and *Caulobacter phage rogue*) in the genus *Phicbklikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.025bB create a new genus in the family *Siphoviridae*

2013.025cB name the new genus *Phicbklikevirus*

2013.025dB designate *Caulobacter phage phicbk* as the type species of the new genus

2013.026a-dB.A.v3.Sap6likevirus [45]

2013.026aB create 5 species (*Enterococcus phage SAP6*, *Enterococcus phage BC611*, *Enterococcus phage IMEEF1*, *Streptococcus phage SPQS1* and *Enterococcus phage VD13*) in the genus *Sap6likevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.026bB create a new genus in the family *Siphoviridae*

2013.026cB name the new genus *Sap6likevirus*

2013.026dB designate *Enterococcus phage SAP6* as the type species of the new genus

2013.027a-dB.A.v3.Tp2unalikevirus [46]

2013.027aB create 2 species, *Bacillus phage TP21* and *Bacillus phage BMBtp2*, in the genus *Tp2unalikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.027bB create a new genus in the family *Siphoviridae*

2013.027cB name the new genus *Tp2unalikevirus*

2013.027dB designate *Bacillus phage TP21* as the type species of the new genus

2013.028a-dB.A.v3.Xp10likevirus [47]

2013.028aB create 5 species (*Xanthomonas phage Xp10*, *Xanthomonas phage OP1*, *Xanthomonas phage Xop411*, *Xanthomonas phage phil7* and *Xanthomonas phage CP1*) in the genus *Xp10likevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.028bB create a new genus in the family *Siphoviridae*

2013.028cB name the new genus *Xp10likevirus*

2013.028dB designate *Xanthomonas phage Xp10* as the type species of the new genus

2013.029a-dB.A.v3.Hk578likevirus [48]

2013.029aB create 5 species (*Escherichia phage HK578*, *Sodalis phage SO-1*, *Shigella phage EP23*, *Enterobacteria phage JL1* and *Enterobacteria phage SSL-2009a*) in the genus *Hk578likevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.029bB create a new genus in the family *Siphoviridae*

2013.029cB name the new genus *Hk578likevirus*

2013.029dB designate *Escherichia phage HK578* as the type species of the new genus

2013.030a-dB.A.v3.Jerseylikevirus [49]

2013.030aB create 6 species (*Salmonella phage jersey*, *Salmonella phage SETP3*, *Salmonella phage Ent1*, *Salmonella phage SE2*, *Salmonella phage wksl3* and *Salmonella phage SS3e*) in the genus *Jerseylikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.030bB create a new genus in the family *Siphoviridae*

2013.030cB name the new genus *Jerseylikevirus*

2013.030dB designate *Salmonella phage jersey* as the type species of the new genus

2013.031a-dB.A.v3.Phie125likevirus [50]

2013.031aB create 3 species (*Burkholderia phage phie125*, *Burkholderia phage phi1026b* and *Burkholderia phage phi6442*) in the genus *Phie125likevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.031bB create a new genus in the family *Siphoviridae*

2013.031cB name the new genus *Phie125likevirus*

2013.031dB designate *Burkholderia phage phie125* as the type species of the new genus

2013.032a-dB.A.v3.Wbetalikevirus [51]

2013.032aB create species *Bacillus phage Wbeta* in the genus *Wbetalikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.032bB create a new genus in the family *Siphoviridae*

2013.032cB name the new genus *Wbetalikevirus*

2013.032dB designate *Bacillus phage Wbeta* as the type species of the new genus

2013.033aB.A.v3.PhiC3unalikeviruses_2sp [52]

2013.033aB create 2 species, *Streptomyces phage phibt1* and *Streptomyces phage TG1*, in the genus *PhiC3unalikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.034a-dB.A.v4.Sfi1unalikevirus [53]

2013.034aB create 5 species (*Streptococcus phage Sfi11*, *Streptococcus phage O1205*, *Streptococcus phage 2972*, *Streptococcus phage Alq132* and *Streptococcus phage 858*) in the genus *Sfi1unalikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.034bB create a new genus in the family *Siphoviridae*

2013.034cB name the new genus *Sfi1unalikevirus*

2013.034dB designate *Streptococcus phage Sfi11* as the type species of the new genus

2013.035a-dB.A.v3.Skunalikevirus [54]

2013.035aB create 17 species (*Lactococcus phage SK1*, *Lactococcus phage bIl170*, *Lactococcus phage jj50*, *Lactococcus phage 712*, *Lactococcus phage P008*, *Lactococcus phage Sl4*, *Lactococcus phage CB13*, *Lactococcus phage CB14*, *Lactococcus phage CB19*, *Lactococcus phage CB20*, *Lactococcus phage Bibb29*, *Lactococcus phage P2*, *Lactococcus phage Ascc273*, *Lactococcus phage Ascc191*, *Lactococcus phage Ascc281*, *Lactococcus phage Ascc532* and *Lactococcus phage Ascc465*) in the genus *Skunalikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.035bB create a new genus in the family *Siphoviridae*

2013.035cB name the new genus *Skunalikevirus*

2013.035dB designate *Lactococcus phage SK1* as the type species of the new genus

2013.036a-dB.A.v3.Sfi21dtunalikevirus [55]

2013.036aB create 5 species (*Streptococcus phage DT1*, *Streptococcus phage Sfi19*, *Streptococcus phage Sfi21*, *Streptococcus phage Abc2 and Streptococcus phage 7201*) in the genus *Sfi21dtunalikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.036bB create a new genus in the family *Siphoviridae*

2013.036cB name the new genus *Sfi21dtunalikevirus*

2013.036dB designate *Streptococcus phage DT1* as the type species of the new genus

2013.037a-dB.A.v3.Andromedalikevirus [56]

2013.037aB create 8 species (*Bacillus phage andromeda*, *Bacillus phage eoghan*, *Bacillus phage taylor*, *Bacillus phage curly*, *Bacillus phage finn*, *Bacillus phage glittering*, *Bacillus phage blastoid* and *Bacillus phage riggi*) in the genus *Andromedalikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.037bB create a new genus in the family *Siphoviridae*

2013.037cB name the new genus *Andromedalikevirus*

2013.037dB designate *Bacillus phage andromeda* as the type species of the new genus

2013.038a-dB.A.v4.Bcep22likevirus [57]

2013.038aB create 4 species (*Burkholderia phage Bcep22*, *Burkholderia phage Bcepil02*, *Burkholderia phage Dc1* and *Burkholderia phage Bcepmigl*) in the genus *Bcep22likevirus* (new), family *Podoviridae*, order *Caudovirales*

2013.038bB create a new genus in the family *Podoviridae*

2013.038cB name the new genus *Bcep22likevirus*

2013.038dB designate *Burkholderia phage Bcep22* as the type species of the new genus

2013.039a-dB.A.v3.Chilikevirus [58]

2013.039aB create 5 species (*Salmonella phage Chi*, *Salmonella phage SPN19*, *Salmonella phage iEPS5*, *Salmonella phage FSLSP088* and *Salmonella phage FSLSP030*) in the genus *Chilikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.039bB create a new genus in the family *Siphoviridae*

2013.039cB name the new genus *Chilikevirus*

2013.039dB designate *Salmonella phage Chi* as the type species of the new genus

2013.040a-dB.A.v4.F116likevirus [59]

2013.040aB   create 2 species, *Pseudomonas phage F116* and *Pseudomonas phage H66*, in the genus *F116likevirus* (new), family *Podoviridae*, order *Caudovirales*

2013.040bB   create a new genus in the family *Podoviridae*

2013.040cB   name the new genus *F116likevirus*

2013.040dB   designate *Pseudomonas phage F116* as the type species of the new genus

2013.042a-lB.A.v3.Siphoviridae_3gen [60, 61]

2013.042aB   create 6 species (*Staphylococcus phage 3a*, *Staphylococcus phage 42e*, *Staphylococcus phage 47*, *Staphylococcus phage Ipla35*, *Staphylococcus phage Slt* and *Staphylococcus phage Phi12*) in the genus *3alikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.042bB   create 3 species (*Staphylococcus phage 77*, *Staphylococcus phage 13* and *Staphylococcus phage Pvl108*) in the genus *77likevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.042cB   create 31 species (*Staphylococcus phage 11*, *Staphylococcus phage 187*, *Staphylococcus phage 29*, *Staphylococcus phage 37*, *Staphylococcus phage 52a*, *Staphylococcus phage 53*, *Staphylococcus phage 55*, *Staphylococcus phage 69*, *Staphylococcus phage 71*, *Staphylococcus phage 80*, *Staphylococcus phage 80alpha*, *Staphylococcus phage 85*, *Staphylococcus phage 88*, *Staphylococcus phage 92*, *Staphylococcus phage 96*, *Staphylococcus phage Cnph82*, *Staphylococcus phage Ew*, *Staphylococcus phage Ph15*, *Staphylococcus phage Phieta*, *Staphylococcus phage Phieta2*, *Staphylococcus phage Phieta3*, *Staphylococcus phage phimr11*, *Staphylococcus phage phimr25*, *Staphylococcus phage Ipla88*, *Staphylococcus phage Sap26*, *Staphylococcus phage X2*, *Staphylococcus phage phinm1*, *Staphylococcus phage phinm4*, *Staphylococcus phage Ipla5*, *Staphylococcus phage Ipla7* and *Staphylococcus phage phinm2*) in the genus *Phietalikevirus* (new), family *Siphoviridae*, order *Caudovirales*

2013.042dB   create a new genus in the family *Siphoviridae*

2013.042eB   name the new genus *3alikevirus*

2013.042fB   designate *Staphylococcus phage 3a* as the type species of the new genus

2013.042gB   create a new genus in the family *Siphoviridae*

2013.042hB   name the new genus *77alikevirus*

2013.042iB   designate *Staphylococcus phage 77* as the type species of the new genus

2013.042jB   create a new genus in the family *Siphoviridae*

2013.042kB   name the new genus *Phietalikevirus*

2013.042lB   designate *Staphylococcus phage Phieta* as the type species of the new genus

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2013.004aV.A.v2.Phlebovirus_sp [62]

2013.004aV   create species *SFTS virus* in the genus *Phlebovirus*, family *Bunyaviridae*

2014.001aV.A.v1. Orthohepadnavirus_sp [63]

2014.001aV   create species *Bat hepatitis virus* in genus *Orthohepadnavirus*, family *Hepadnaviridae*

2014.002aV.A.v2.Rotavirus_3sp [64]

2014.002aV   create 3 species (*Rotavirus F*, *Rotavirus G*, and *Rotavirus H*), in the genus *Rotavirus*, subfamily *Sedoreovirinae*, family *Reoviridae*

2014.004aV.A.v2.Tospovirus_sp [65]

2014.004aV   create species *Iris yellow spot virus* in the genus *Tospovirus*, family *Bunyaviridae*

2014.005aV.A.v2.Tospovirus_sp [66]

2014.005aV   create species *Watermelon bud necrosis virus* in the genus *Tospovirus*, family *Bunyaviridae*

2014.007aV.A.v2.Nyavirus_sp [67]

2014.007aV   create species *Sierra Nevada nyavirus* in the genus *Nyavirus*, family *Nyamiviridae*, order *Mononegavirales*

2014.008a-hV.A.v6.Hepeviridae [68]

2014.008aV   create species *Piscihepevirus A* in the genus *Piscihepevirus* (new), family *Hepeviridae*

2014.008bV   create 2 species (*Orthohepevirus C* and *Orthohepevirus D*) in the genus *Hepevirus* (proposed name, *Orthohepevirus*) family *Hepeviridae*

2014.008cV   create a new genus in the family *Hepeviridae*

2014.008dV   name the genus *Piscihepevirus*

2014.008eV   designate *Piscihepevirus A* as type species of the new genus

2014.008fV   remove species *Avian hepatitis E virus* from its unassigned position within the family *Hepeviridae*

2014.008gV   reassign the above species to the genus *Hepevirus* (proposed name, *Orthohepevirus*)

2014.008hV   (i) Change the name of the genus *Hepevirus* to *Orthohepevirus*. (ii) Change the name of the species *Hepatitis E virus* to *Orthohepevirus A*. (iii) Change the name of the species *Avian hepatitis E virus* to *Orthohepevirus B*.

2014.010a,bV.A.v2.Bornaviridae_spren [69, 70]

2014.010aV   create 4 species (*Psittaciform 1 bornavirus*, *Passeriform 1 bornavirus*, *Passeriform 2 bornavirus* and *Waterbird 1 bornavirus*) in the genus *Bornavirus*, family *Bornaviridae*

2014.010bV   In the genus *Bornavirus*, change the name of species *Borna disease virus* to *Mammalian 1 bornavirus*

2014.011a-dV.A.v2.Reptarenavirus [71]

2014.011aV   create 3 species (*Alethinophid 1 reptarenavirus*, *Alethinophid 2 reptarenavirus* and *Alethinophid 3 reptarenavirus*) in the genus *Reptarenavirus* (new), family *Arenaviridae*

2014.011bV   create a new genus in the family *Arenaviridae*

2014.011cV   name the genus *Reptarenavirus*

2014.011dV   designate *Alethinophid 1 reptarenavirus* as type species of the new genus

2014.012aV.A.v3.Arenavirus_ren [72]

2014.012aV   in the family *Arenaviridae*, change the name of genus *Arenavirus* to *Mammarenavirus* and convert the names of its constituent species to a binomial format, by changing the word *virus* to *mammarenavirus* in each existing name, [example: *Allpahuayo virus* (current name) changed to *Allpahuayo mammarenavirus* (new name)]

2014.013aV.A.v3. Mammarenavirus_2sp [73]

2014.013aV   create species *Lunk mammarenavirus* and *Merino Walk mammarenavirus* in the genus *Arenavirus* (proposed name *Mammarenavirus*), family *Arenaviridae*

2014.014aV.A.v2.Cardiovirus_sp [74]

2014.014aV   create species *Cardiovirus C* within the genus *Cardiovirus*, family *Picornaviridae*

2014.015a-dV.A.v3.Kunsagivirus [75]

2014.015aV   create species *Kunsagivirus A* in genus *Kunsagivirus* (new), family *Picornaviridae*

2014.015bV   create a new genus in the family *Picornaviridae*

2014.015cV   name the genus *Kunsagivirus*

2014.015dV   designate *Kunsagivirus A* as type species of the new genus

2014.016aV.A.v1.Picornaviridae_spren [76]

2014.016aV   rename 12 species in the family *Picornaviridae* as follows:

Genus	**Old name**	**New name**
Avihepatovirus	*Duck hepatitis A virus*	*Avihepatovirus A*
Cardiovirus	*Encephalomyocarditis virus*	*Cardiovirus A*
Cardiovirus	*Theilovirus*	*Cardiovirus B*
Erbovirus	*Equine rhinitis B virus*	*Erbovirus A*
Hepatovirus	*Hepatitis A virus*	*Hepatovirus A*
Parechovirus	*Human parechovirus*	*Parechovirus A*
Parechovirus	*Ljungan virus*	*Parechovirus B*
Sapelovirus	*Porcine sapelovirus*	*Sapelovirus A*
Sapelovirus	*Simian sapelovirus*	*Sapelovirus B*
Senecavirus	*Seneca Valley virus*	*Senecavirus A*
Teschovirus	*Porcine teschovirus*	*Teschovirus A*
Tremovirus	*Avian encephalomyelitis virus*	*Tremovirus A*

2014.017a-dV.A.v3.Sakobuvirus [77]

2014.017aV   create species *Sakobuvirus A* in genus *Sakobuvirus* (new), family *Picornaviridae*

2014.017bV   create a new genus in the family *Picornaviridae*

2014.017cV   name the genus *Sakobuvirus*

2014.017dV   designate *Sakobuvirus A* as type species of the new genus

2014.018a-dV.A.v3.Sicinivirus [78]

2014.018aV   create species *Sicinivirus A* in genus *Sicinivirus* (new), family *Picornaviridae*

2014.018bV   create a new genus in the family *Picornaviridae*

2014.018cV   name the genus *Sicinivirus*

2014.018dV   designate *Sicinivirus A* as type species of the new genus

All of the above taxonomic proposals were approved by the members who voted, mostly unanimously and always >96 %. The return rate of votes was approximately 69 %. These taxonomic changes are now part of the official ICTV taxonomy and represent changes to the taxonomy published in the Ninth Report [79] and are in addition to changes approved in February 2012 [80], February 2013 [81] and March 2014 [82]. An up-to-date list of all approved taxa can be found on the ICTV online web site: http://www.ictvonline.org/msl.htm.

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