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Ratification vote on taxonomic proposals to the International Committee on Taxonomy of Viruses (2016)

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Abstract This article lists the changes to virus taxonomy approved and ratified by the International Committee on Taxonomy of Viruses (ICTV) in April 2016.

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 (http://www.ictvonline.org/statutes.asp), proposals submitted to the ICTV Executive Committee

(EC) undergo a review process that involves input from the ICTV Study Groups (SGs) and Subcommittees (SCs), other interested virologists, and the EC. After final approval by the EC, proposals are then presented for ratification to the full ICTV membership by publication on an ICTV web site (http://www.ictvonline.org/) followed by an electronic vote. The latest set of proposals approved by the EC was made available on the ICTV website by January 2016 (https://talk.ictvonline.org/files/proposals/). A list of these

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proposals was then emailed on 28 March 2016 to the 148 members of ICTV, namely the EC Members, Life Members, ICTV Subcommittee Members (including the SG chairs) and ICTV National Representatives. Members were then requested to vote on whether to ratify the taxonomic proposals (voting closed on 29 April 2016).

This report summarises the taxonomic proposals that were ratified by ICTV members. Note that in accordance with ICTV convention, all approved taxon names are italicized. The proposals are listed by ICTV Subcommittee, and then the filename as posted on-line. Each proposal is formally cited so that a list of the authors appears in the References. This provides a means of acknowledging the authors' efforts and also links to the specific proposal on the ICTV website. These documents remain available for any who wish to see details of the viruses assigned to the new taxa or any other details of the proposals.

Bacterial and Archaeal Viruses (code B)
2015.001a-dB.A.v2.Cba41virus [1]

2015.001aB	create 2 species in genus *Cba41virus* (new) in the family *Podoviridae*: *Cellulophaga virus Cba41*, *Cellulophaga virus Cba172*
2015.001bB	create a new genus in the family *Podoviridae*, order *Caudovirales*
2015.001cB	name the genus *Cba41virus*
2015.001dB	designate *Cellulophaga virus Cba41* as type species of the new genus

2015.002a-dB.A.v2.Cba181virus [2]

2015.002aB	create 3 species in genus *Cba181virus* (new) in the family *Siphoviridae*: *Cellulophaga virus Cba181*, *Cellulophaga virus Cba121*, *Cellulophaga virus Cba171*
2015.002bB	create a new genus in family *Siphoviridae*, order *Caudovirales*
2015.002cB	name the genus *Cba181virus*
2015.002dB	designate *Cellulophaga virus Cba181* as type species of the new genus

2015.003a-dB.A.v2.Cbastvirus [3]

2015.003aB	create species in genus *Cbastvirus* (new) in the family *Siphoviridae*: *Cellulophaga virus ST*
2015.003bB	create a new genus in the family *Siphoviridae*, order *Caudovirales*
2015.003cB	name the genus *Cbastvirus*
2015.003dB	designate *Cellulophaga virus ST* as type species of the new genus

2015.004a-dB.A.v2.Ff47virus [4]

2015.004aB	create 2 species in genus *Ff47virus* (new) in the family *Siphoviridae*: *Mycobacterium virus Ff47*, *Mycobacterium virus Muddy*

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2015.004bB   create a new genus in the family
              Siphoviridae, order *Caudovirales*
2015.004cB   name the genus *Ff47virus*
2015.004dB   designate *Mycobacterium virus Ff47* as type
              species of the new genus

2015.005a-gB.A.v3.Kayvirus [5]

2015.005aB   create 6 species in genus *Kayvirus* (new) in
              the subfamily *Spounavirinae*, family
              Myoviridae: *Staphylococcus virus JD7*,
              Staphylococcus virus SA12, *Staphylococcus
              virus P108*, *Staphylococcus virus G15*,
              Staphylococcus virus MCE2014,
              Staphylococcus virus S253
2015.005bB   create a new genus in the subfamily
              Spounavirinae, family *Myoviridae*, order
              *Caudovirale*s
2015.005cB   name the genus *Kayvirus*
2015.005dB   designate *Staphylococcus virus K* as type
              species of the new genus
2015.005eB   remove species *Staphylococcus phage G1*
              and *Staphylococcus phage K* from genus
              Twortlikevirus
2015.005fB   reassign the above two species to genus
              Kayvirus
2015.005gB   change the name of species *Staphylococcus
              phage G1* and *Staphylococcus phage K* to
              Staphylococcus virus G1 and
              Staphylococcus virus K, respectively

2015.006aB.A.v2.Phage_Genera_ren [6]

2015.006aB   change the names of all 79 existing phage
              genera by deleting *like* from each name (e.g.
              Andromedalikevirus to *Andromedavirus*).
              Also make the following individual name
              changes (proposed name in brackets):
              *3alikevirus (Triavirus), 77likevirus
              (Biseptimavirus), Ahjdlikevirus (P68virus),
              Bppunalikevirus (Bpp1virus),
              Cjwunalikevirus (Cjw1virus),
              Cp8unalikevirus (Cp8virus),
              Felixounalikevirus (Felixo1virus),
              Halolikevirus (Liefievirus), Hapunalikevirus
              (Hapunavirus), Hpunalikevirus (Hp1virus),
              I3likevirus (Bxz1virus), Iebhlikevirus
              (Cecivirus), Pbiunalikevirus (Pbi1virus),
              Pbunalikevirus (Pbunavirus),
              Pgonelikevirus (Pg1virus),
              Phic3unalikevirus (Phic31virus),
              Phicd119likevirus (Cd119virus),
              Phie125likevirus (E125virus), Phihlikevirus*

*(Myohalovirus), Phijlunalikevirus
(Phijl1virus), Psimunalikevirus
(Psimunavirus), Punalikevirus (P1virus),
Sfi1unalikevirus (Sfi11virus),
Sfi21dtunalikevirus (Sfi21dt1virus),
Skunalikevirus (Sk1virus), Spounalikevirus
(Spo1virus), Tp2unalikevirus (Tp21virus),
Tunalikevirus (T1virus), Viunalikevirus
(Vi1virus)*

2015.007a-dB.A.v1.Kp34virus [7]

2015.007aB   create 5 species in genus *Kp34virus* (new)
              in the family *Podoviridae*: *Klebsiella virus
              KP34, Klebsiella virus SU503, Klebsiella
              virus SU552A, Klebsiella virus F19,
              Klebsiella virus K244*
2015.007bB   create a new genus in the family
              Podoviridae, order *Caudovirales*
2015.007cB   name the genus *Kp34virus*
2015.007dB   designate *Klebsiella virus KP34* as type
              species of the new genus

2015.008a-dB.A.v3.Lit1virus [8]

2015.008aB   create 3 species in genus *Lit1virus* (new) in
              the family *Podoviridae*: *Pseudomonas virus
              LIT1, Pseudomonas virus PA26,
              Pseudomonas virus Ab09*
2015.008bB   create a new genus in the family
              Podoviridae, order *Caudovirales*
2015.008cB   name the genus *Lit1virus*
2015.008dB   designate *Pseudomonas virus LIT1* as type
              species of the new genus

2015.009aB.A.v3.Luz24virus_5sp [9]

2015.009aB   create 5 species in genus *Luz24virus* in the
              family *Podoviridae*: *Pseudomonas virus
              CHU, Pseudomonas virus Ab22,
              Pseudomonas virus PAA2, Pseudomonas
              virus TL, Pseudomonas virus PaP4*

2015.010a-dB.A.v2.Nonanavirus [10]

2015.010aB   create 2 species in genus *Nonanavirus*
              (new) in the family *Podoviridae*:
              *Salmonella virus 9NA, Salmonella virus
              SP069*
2015.010bB   create a new genus in the family
              Podoviridae, order *Caudovirales*
2015.010cB   name the genus *Nonanavirus*

2015.010dB	designate *Salmonella virus 9NA* as type species of the new genus

2015.011a-dB.A.v2.P70virus [11]

2015.011aB	create 5 species in genus *P70virus* (new) in the family *Siphoviridae*: *Listeria virus P70, Listeria virus LP26, Listeria virus LP110, Listeria virus LP114, Listeria virus LP37*
2015.011bB	create a new genus in the family *Siphoviridae*, order *Caudovirales*
2015.011cB	name the genus *P70virus*
2015.011dB	designate *Listeria virus P70* as type species of the new genus

2015.012a-eB.A.v3.P100virus [12]

2015.012aB	create a new genus in the subfamily *Spounavirinae*, family *Myoviridae*, order *Caudovirales*
2015.012bB	name the genus *P100virus*
2015.012cB	designate *Listeria virus P100* as type species of the new genus
2015.012dB	remove species *Listera phage P100* and *Listera phage A511* from genus *Twortlikevirus*
2015.012eB	reassign the above two species to genus *P100virus*

2015.013a-cB.A.v3.Biquartavirus [13]

2015.013aB	create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.013bB	name the genus *Biquartavirus*
2015.013cB	designate *Aeromonas virus 44RR2* (see 2015.020aeB and 2015.025aB) as type species of the new genus

2015.014aB.A.v3.Phieco32virus_4sp [14]

2015.014aB	create 4 species within the genus *Phieco32virus,* family *Podoviridae*: *Escherichia virus SU10, Escherichia virus NJ01, Escherichia virus ECB2, Escherichia virus Septima11*

2015.015a-dB.A.v2.Psavirus [15]

2015.015aB	create 2 species in genus *Psavirus* (new) in the family *Siphoviridae*: *Listeria virus PSA, Listeria virus LP302*
2015.015bB	create a new genus in the family *Siphoviridae*, order *Caudovirales*

2015.015cB	name the genus *Psavirus*
2015.015dB	designate *Listeria virus PSA* as type species of the new genus

2015.016a-dB.A.v3.Secunda5virus [16]

2015.016aB	create 4 species in genus *Secunda5virus* (new), family *Myoviridae*: *Aeromonas virus AS4, Stenotrophomonas virus IME13, Aeromonas virus Aes12, Aeromonas virus Aes508*
2015.016bB	create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.016cB	name the genus *Secunda5virus*
2015.016dB	designate *Aeromonas virus 25* (see 2015.020adB) as type species of the new genus

2015.017a-dB.A.v3.Sextaecvirus [17]

2015.017aB	create 2 species in genus *Sextaecvirus* (new) in the family *Siphoviridae*: *Staphylococcus virus Sextaec, Staphylococcus virus SEP9*
2015.017bB	create a new genus in the family *Siphoviridae*, order *Caudovirales*
2015.017cB	name the genus *Sextaecvirus*
2015.017dB	designate *Staphylococcus virus Sextaec* as type species of the new genus

2015.018a-dB.A.v2.Silviavirus [18]

2015.018aB	create 2 species in genus *Silviavirus* (new), family *Myoviridae*: *Staphylococcus virus Remus, Staphylococcus virus SA11*
2015.018bB	create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.018cB	name the genus *Silviavirus*
2015.018dB	designate *Staphylococcus virus Remus* as type species of the new genus

2015.019a-abB.A.v3.Tunavirinae [19]

2015.019aB	create 2 species in genus *Tunalikevirus* (proposed name *T1virus*) in the family *Siphoviridae*
2015.019bB	create 2 species in genus *Tlsvirus* (new) in the subfamily *Tunavirinae* (new), family *Siphoviridae*: *Escherichia virus ADB2, Shigella virus PSf2, Salmonella virus SP126, Citrobacter virus Stevie*
2015.019cB	create a new genus in the subfamily *Tunavirinae* (new), family *Siphoviridae*, order *Caudovirales*
2015.019dB	name the genus *Tlsvirus*

2015.019eB	designate *Escherichia phage Tls* (proposed name *Escherichia virus Tls*) as type species of the new genus
2015.019fB	create species in genus *Rtpvirus* (new) in the family *Siphoviridae*: *Escherichia virus ACGM12*
2015.019gB	create a new genus in the subfamily *Tunavirinae* (new), family *Siphoviridae*, order *Caudovirales*
2015.019hB	name the genus *Rtpvirus*
2015.019iB	designate *Escherichia phage Rtp* (proposed name *Escherichia virus RTP*) as type species of the new genus
2015.019jB	create 2 species in genus *Kp36virus* (new) in the subfamily *Tunavirinae* (new), family *Siphoviridae*: *Klebsiella virus KP36, Klebsiella virus 1513*
2015.019kB	create a new genus in the subfamily *Tunavirinae* (new), family *Siphoviridae*, order *Caudovirales*
2015.019lB	name the genus *Kp36virus*
2015.019mB	designate *Klebsiella virus KP36* as type species of the new genus
2015.019nB	create 5 species in genus *Rogue1virus* (new) in the subfamily *Tunavirinae* (new), family *Siphoviridae*: *Escherichia virus AHS24, Escherichia virus KP26, Escherichia virus AHP42, Escherichia virus AKS96, Escherichia virus E41c*
2015.019oB	create a new genus in the subfamily *Tunavirinae* (new), family *Siphoviridae*, order *Caudovirales*
2015.019pB	name the genus *Rogue1virus*
2015.019qB	designate *Escherichia phage Rogue1* (proposed name *Escherichia virus Rogue1*) as type species of the new genus
2015.019rB	create a new subfamily in the family *Siphoviridae*, order *Caudovirales*
2015.019sB	name the new subfamily *Tunavirinae*
2015.019tB	assign to the new subfamily the 5 genera, *Tunalikevirus* (existing, proposed name *T1virus*), *Rtpvirus* (new), *Tlsvirus* (new), *Kp36virus* (new) and *Rogue1virus* (new)
2015.019uB	remove genus *Tunalikevirus* from its position of unassigned in the family *Siphoviridae*
2015.019vB	reassign the above genus to the subfamily *Tunavirinae* (new), family *Siphoviridae*
2015.019wB	remove 7 species (*Cronobacter phage Esp2949-1, Enterobacter phage F20, Escherichia phage Eb49, Escherichia phage Jk06, Escherichia phage Rogue1, Escherichia phage Rtp* and *Escherichia phage Tls*) from genus *Tunalikevirus*
2015.019xB	reassign species *Cronobacter phage Esp2949-1* to be unassigned in the subfamily *Tunavirinae* (new)
2015.019yB	reassign species *Escherichia phage Tls* (proposed name *Escherichia virus Tls*) to the new genus *Tlsvirus* in the subfamily *Tunavirinae* (new)
2015.019zB	reassign species *Enterobacter phage F20* (proposed name *Enterobacter virus F20*) to the new genus *Kp36virus* in the subfamily *Tunavirinae* (new)
2015.019aaB	reassign species *Escherichia phage Rtp* (proposed name *Escherichia virus RTP*) to the new genus *Rtpvirus* in the subfamily *Tunavirinae* (new)
2015.019abB	reassign species *Escherichia virus Jk06* (proposed name *Escherichia virus Jk06*) and *Escherichia phage Eb49* (proposed name *Escherichia phage Eb49*) and *Escherichia phage Rogue1* (proposed name *Escherichia virus Rogue1*) to the new genus *Rogue1virus* in the subfamily *Tunavirinae* (new)

2015.020a-aeB.A.v4.Tevenvirinae [20]

2015.020aB	create 10 species in genus *T4likevirus* (proposed name *T4virus*) in the subfamily *Tevenvirinae* (new), family *Myoviridae*: *Escherichia virus AR1, Escherichia virus Ime09, Escherichia virus E112, Escherichia virus ECML134, Escherichia virus RB3, Escherichia virus C40, Shigella virus Shfl2, Shigella virus Pss1, Yersinia virus PST, Yersinia virus D1*
2015.020bB	create a new genus in the subfamily *Tevenvirinae* (new), family *Myoviridae*, order *Caudovirales*
2015.020cB	name the genus *Rb49virus*
2015.020dB	designate *Escherichia phage RB49* (existing species) to be renamed *Escherichia virus RB49* as type species of the new genus
2015.020eB	create 3 species in genus *Rb69virus* (new) in the subfamily *Tevenvirinae* (new), family *Myoviridae*: *Escherichia virus JS09, Escherichia virus HX01, Shigella virus UTAM*
2015.020fB	create a new genus in the subfamily *Tevenvirinae* (new), family *Myoviridae*, order *Caudovirales*

2015.020gB	name the genus *Rb69virus*
2015.020hB	designate *Escherichia phage RB69* (existing species) to be renamed *Escherichia virus RB69* as type species of the new genus
2015.020iB	create 4 species in genus *Js98virus* (new) in the subfamily *Tevenvirinae* (new), family *Myoviridae*: *Escherichia virus IME08*, *Escherichia virus Bp7*, *Escherichia virus JS10*, *Escherichia virus VR5*
2015.020jB	create a new genus in the subfamily *Tevenvirinae* (new), family *Myoviridae*, order *Caudovirales*
2015.020kB	name the genus *Js98virus*
2015.020lB	designate *Escherichia phage JS98* (existing species) to be renamed *Escherichia virus JS98* as type species of the new genus
2015.020mB	create 5 species in genus *Sp18virus* (new) in the subfamily *Tevenvirinae* (new), family *Myoviridae*: *Shigella virus SP18*, *Escherichia virus VR7*, *Escherichia virus VR20*, *Escherichia virus VR25*, *Escherichia virus VR26*
2015.020nB	create a new genus in the subfamily *Tevenvirinae* (new), family *Myoviridae*, order *Caudovirales*
2015.020oB	name the genus *Sp18virus*
2015.020pB	designate *Shigella virus SP18* as type species of the new genus
2015.020qB	create 2 species in genus *Sp16virus* (new) in the subfamily *Tevenvirinae* (new), family *Myoviridae*: *Salmonella virus STML198*, *Salmonella virus S16*
2015.020rB	create a new genus in the subfamily *Tevenvirinae* (new), family *Myoviridae*, order *Caudovirales*
2015.020sB	name the genus *Sp16virus*
2015.020tB	designate *Salmonella virus S16* as type species of the new genus
2015.020uB	create 2 species in genus *Cc31virus* (new) in the subfamily *Tevenvirinae* (new), family *Myoviridae*: *Enterobacter virus CC31*, *Enterobacter virus PG7*
2015.020vB	create a new genus in the subfamily *Tevenvirinae* (new), family *Myoviridae*, order *Caudovirales*
2015.020wB	name the genus *Cc31virus*
2015.020xB	designate *Enterobacter virus CC31* as type species of the new genus
2015.020yB	remove 12 species (*Aeromonas phage 25*, *Aeromonas phage 31*, *Aeromonas phage 44RR2.8t*, *Enterobacteria phage SV14*,

	Escherichia phage JS98, *Escherichia phage phi1*, *Escherichia phage RB16*, *Escherichia phage RB32*, *Escherichia phage RB43*, *Escherichia phage RB49*, *Escherichia phage RB69* and *Pseudomonas phage 42*) from genus *T4likevirus*
2015.020zB	reassign 2 of the above species (*Escherichia phage RB49* [proposed name *Escherichia virus RB49*] and *Escherichia phage phi1* [proposed name *Escherichia virus phi1*]) to genus *Rb49virus* (new)
2015.020aaB	reassign *Escherichia phage RB69* (proposed name *Escherichia virus RB69*) to genus *Rb69virus* (new)
2015.020abB	reassign *Escherichia phage JS98* (proposed name *Escherichia virus JS98*) to genus *JS98virus* (new)
2015.020acB	reassign 5 of the above species (*Enterobacteria phage SV14* [proposed name *Enterobacteria virus SV14*], *Escherichia phage Rb16* [proposed name *Escherichia virus Rb16*], *Escherichia phage RB32* [proposed name *Escherichia virus RB32*], *Escherichia phage Rb43* [proposed name *Escherichia virus Rb43*] and *Pseudomonas phage 42* [proposed name *Pseudomonas virus 42*]) to be unassigned in subfamily *Tevenvirinae*
2015.020adB	reassign *Aeromonas phage 25* (proposed name *Aeromonas virus 25*) and *Aeromonas phage 31* (proposed name *Aeromonas virus 31*) to genus *Secunda5virus* (new)
2015.020aeB	reassign *Aeromonas phage 44RR2.8t* (proposed name *Aeromonas virus 44RR2*) to genus *Biquartavirus* (new)

2015.021a,bB.A.v3.T5virus_4sp1rem [21]

2015.021aB	create 4 species in genus *T5likevirus* (proposed name *T5virus*), family *Siphoviridae*: *Escherichia virus FFH1*, *Salmonella virus Shivani*, *Salmonella virus Stitch*, *Escherichia virus DT57C*
2015.021bB	Remove (abolish) species *Vibrio phage 149* (type IV) from genus *T5likevirus*

2015.022a-dB.A.v2.Tg1virus [22]

2015.022aB	create 2 species in genus *Tg1virus* (new) in the family *Myoviridae*: *Yersinia virus TG1*, *Yersinia virus R1RT*
2015.022bB	create a new genus in the family *Myoviridae*, order *Caudovirales*

2015.022cB	name the genus *Tg1virus*
2015.022dB	designate *Yersinia virus TG1* as type species of the new genus

2015.023a-oB.A.v2.Vequintavirinae [23]

2015.023aB	create 4 species in genus *V5virus* (new) in the subfamily *Vequintavirinae* (new), family *Myoviridae*: *Escherichia virus V5, Escherichia virus JES2013, Escherichia virus FV3, Escherichia virus FFH2*
2015.023bB	create 4 species in genus *Se1virus* (new) in the subfamily *Vequintavirinae* (new), family *Myoviridae*: *Salmonella virus SE1, Salmonella virus SSE121, Escherichia virus 4MG, Cronobacter virus GAP31*
2015.023cB	create 3 species in genus *Cr3virus* (new) in the subfamily *Vequintavirinae* (new), family *Myoviridae*:, *Cronobacter virus CR3, Cronobacter virus CR8, Cronobacter virus CR9*
2015.023dB	create a new genus in the subfamily *Vequintavirinae* (new), family *Myoviridae*, order *Caudovirales*
2015.023eB	name the genus *V5virus*
2015.023fB	designate *Escherichia virus V5* as type species of the new genus
2015.023gB	create a new genus in the subfamily *Vequintavirinae* (new), family *Myoviridae*, order *Caudovirales*
2015.023hB	name the genus *Se1virus*
2015.023iB	designate *Salmonella virus SE1* as type species of the new genus
2015.023jB	create a new genus in the subfamily *Vequintavirinae* (new), family *Myoviridae*, order *Caudovirales*
2015.023kB	name the genus *Cr3virus*
2015.023lB	designate *Cronobacter virus CR3* as type species of the new genus
2015.023mB	create a new subfamily in the family *Myoviridae*, order *Caudovirales*
2015.023nB	name the subfamily *Vequintavirinae*
2015.023oB	Assign the following 3 genera to the new subfamily: *V5virus* (new), *Se1virus* (new) and *Cr3virus* (new)

2015.024a-dB.A.v2.Vhmlvirus [24]

2015.024ab	create 3 species in genus *Vhmlvirus* (new) in the family *Myoviridae*: *Vibrio virus VHML, Vibrio virus MAR, Vibrio virus VP585*
2015.024bB	create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.024cB	name the genus *Vhmlvirus*
2015.024dB	designate *Vibrio virus VHML* as type species of the new genus

2015.025aB.A.v4.phagesp_ren [25]

2015.025aB	rename all (522) existing bacterial virus species and 2 archaeal virus species (please refer to the proposal document for a complete list of changes)

2015.026a-rB.A.v3.Bullavirinae [26]

2015.026aB	create a new genus in the subfamily *Bullavirinae* (new), family *Microviridae*
2015.026bB	name the genus *Phix174microvirus*
2015.026cB	designate *Escherichia phage phiX174* (proposed name *Escherichia virus phiX174*) as type species of the new genus
2015.026dB	create 7 species in genus *Alpha3microvirus* (new), subfamily *Bullavirinae* (new), family *Microviridae*: *Escherichia virus WA45, Escherichia virus NC35, Escherichia virus ID21, Escherichia virus NC28, Escherichia virus NC29, Escherichia virus ID32, Escherichia virus ID62*
2015.026eB	create a new genus in the subfamily *Bullavirinae* (new), family *Microviridae*
2015.026fB	name the genus *Alpha3microvirus*
2015.026gB	designate *Escherichia phage alpha3* (proposed name *Escherichia virus alpha3*) as type species of the new genus
2015.026hB	create 2 species in genus *G4microvirus* (new), subfamily *Bullavirinae* (new), family *Microviridae*: *Escherichia virus ID52, Escherichia virus Talmos*
2015.026iB	create a new genus in the subfamily *Bullavirinae* (new), family *Microviridae*
2015.026jB	name the genus *G4microvirus*
2015.026kB	designate *Escherichia phage G4* (proposed name *Escherichia virus G4*) as type species of the new genus
2015.026lB	create a new subfamily in the family *Microviridae*
2015.026mB	name the new subfamily *Bullavirinae*
2015.026nB	assign the following 3 genera to the new subfamily: *Phix174microvirus* (new), *Alpha3microvirus* (new) and *G4microvirus* (new)
2015.026oB	remove 5 species (*Enterobacteria phage alpha3, Enterobacteria phage G4, Enterobacteria phage phiK, Enterobacteria phage phiX174* and *Enterobacteria phage St-1*) from the genus *Microvirus*, family *Microviridae*

2015.026pB   reassign species *Enterobacteria phage phiX174* (proposed name *Enterobacteria virus phiX174*) to the genus *Phix174microvirus* (new), family *Microviridae*

2015.026qB   reassign species *Enterobacteria phage G4* (proposed name *Enterobacteria virus G4*) to the genus *G4microvirus* (new), family *Microviridae*

2015.026rB   reassign 3 species (*Enterobacteria phage alpha3* [proposed name *Escherichia virus alpha3*], *Enterobacteria phage phiK* [proposed name *Escherichia virus phiK*] and *Enterobacteria phage St-1* [proposed name *Escherichia virus St1*]) to the genus *Alpha3microvirus* (new), family *Microviridae*

2015.027aB.A.v3.Viunalikevirus_7sp [27]

2015.027aB   create 7 species in genus *Viunalikevirus* (proposed name *Vi1virus*), family *Myoviridae*, order *Caudovirales*: *Escherichia virus ECML4, Salmonella virus Marshall, Salmonella virus STML131, Salmonella virus SJ3, Salmonella virus Maynard, Salmonella virus Det7, Salmonella virus SJ2*

2015.028a-dB.A.v2.Kpp10virus [28]

2015.028aB   create 3 species in genus *Kpp10virus* (new), family *Myoviridae*: *Pseudomonas virus KPP10, Pseudomonas virus PAKP3, Pseudomonas virus Ab03*

2015.028bB   create a new genus in the family *Myoviridae*, order *Caudovirales*

2015.028cB   name the genus *Kpp10virus*

2015.028dB   designate *Pseudomonas virus KPP10* as type species of the new genus

2015.029a-dB.A.v2.Pakpunavirus [29]

2015.029aB   create 6 species in genus *Pakpunavirus* (new), family *Myoviridae*: *Pseudomonas virus PAKP1, Pseudomonas virus PAKP4, Pseudomonas virus JG004, Pseudomonas virus CAb02, Pseudomonas virus PaP1, Pseudomonas virus CAb1*

2015.029bB   create a new genus in the family *Myoviridae*, order *Caudovirales*

2015.029cB   name the genus *Pakpunavirus*

2015.029dB   designate *Pseudomonas virus PAKP1* as type species of the new genus

2015.030aB.A.v3.Pbunalikevirus_5sp [30]

2015.030aB   create 5 species in genus *Pbunalikevirus* (to be renamed *Pbunavirus*), family *Myoviridae*: *Pseudomonas virus DL60, Pseudomonas virus DL68, Pseudomonas virus JG024, Pseudomonas virus KPP12, Pseudomonas virus Ab28*

2015.031a-dB.A.v1.Vp5virus [31]

2015.031aB   create 3 species in genus *Vp5virus* (new) in the family *Podoviridae*: *Vibrio virus VP5, Vibrio virus VC8, Vibrio virus VP2*

2015.031bB   create a new genus in the family *Podoviridae*, order *Caudovirales*

2015.031cB   name the genus *Vp5virus*

2015.031dB   designate *Vibrio virus VP5* as type species of the new genus

2015.032a-dB.A.v2.Agatevirus [32]

2015.032aB   create 3 species in genus *Agatevirus* (new) in the family *Myoviridae*: *Bacillus virus Agate, Bacillus virus Bobb, Bacillus virus Bp8pC*

2015.032bB   create a new genus in the family *Myoviridae*, order *Caudovirales*

2015.032cB   name the genus *Agatevirus*

2015.032dB   designate *Bacillus virus Agate* as type species of the new genus

2015.033a-dB.A.v2.B4virus [33]

2015.033aB   create 5 species in genus *B4virus* (new) in the family *Myoviridae*: *Bacillus virus B4, Bacillus virus Spock, Bacillus virus Bigbertha, Bacillus virus Troll, Bacillus virus Riley*

2015.033bB   create a new genus in the family *Myoviridae*, order *Caudovirales*

2015.033cB   name the genus *B4virus*

2015.033dB   designate *Bacillus virus B4* as type species of the new genus

2015.034a-dB.A.v2.Bastillevirus [34]

2015.034aB   create 2 species in genus *Bastillevirus* (new) in the family *Myoviridae*: *Bacillus virus Bastille, Bacillus virus CAM003*

2015.034bB   create a new genus in the family *Myoviridae*, order *Caudovirales*

2015.034cB   name the genus *Bastillevirus*

2015.034dB   designate *Bacillus virus Bastille* as type species of the new genus

2015.035a-dB.A.v2.Bc431virus [35]

2015.035aB	create 4 species in genus *Bc431virus* (new) in the family *Myoviridae*: *Bacillus virus Bcp1, Bacillus virus Bc431, Bacillus virus JBP901, Bacillus virus BCP82*
2015.035bB	create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.035cB	name the genus *Bc431virus*
2015.035dB	designate *Bacillus virus Bc431* as type species of the new genus

2015.036a-dD.A.v2.Cp51virus [36]

2015.036aB	create 3 species in genus *Cp51virus* (new) in the family *Myoviridae*: *Bacillus virus CP51, Bacillus virus Shanette, Bacillus virus JL*
2015.036bB	create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.036cB	name the genus *Cp51virus*
2015.036dB	designate *Bacillus virus CP51* as type species of the new genus

2015.037a-dB.A.v3.Sitaravirus [37]

2015.037aB	create 5 species in genus *Sitaravirus* (new) in the family *Siphoviridae*: *Paenibacillus virus Diva, Paenibacillus virus Shelly, Paenibacillus virus Sitara, Paenibacillus virus Rani, Paenibacillus virus Hb10c2*
2015.037bB	create a new genus in the family *Siphoviridae*, order *Caudovirales*
2015.037cB	name the genus *Sitaravirus*
2015.037dB	designate *Paenibacillus virus Diva* as type species of the new genus

2015.038a-dB.A.v3.Nit1virus [38]

2015.038aB	create 3 species in genus *Nit1virus* (new) in the family *Myoviridae*: *Bacillus virus NIT1, Bacillus virus Grass, Bacillus virus SPG24*
2015.038bB	create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.038cB	name the genus *Nit1virus*
2015.038dB	designate *Bacillus virus NIT1* as type species of the new genus

2015.039a-dB.A.v2.Pagevirus [39]

2015.039aB	create 5 species in genus *Pagevirus* (new) in the family *Podoviridae*: *Bacillus virus Page, Bacillus virus Pony, Bacillus virus Pookie, Bacillus virus Palmer, Bacillus virus Pascal*
2015.039bB	create a new genus in the family *Podoviridae*, order *Caudovirales*

2015.039cB	name the genus *Pagevirus*
2015.039dB	designate *Bacillus virus Page* as type species of the new genus

2015.040a-dB.A.v2.Slashvirus [40]

2015.040aB	create 4 species in genus *Slashvirus* (new) in the family *Siphoviridae*: *Bacillus virus Slash, Bacillus virus Staley, Bacillus virus Stahl, Bacillus virus Stills*
2015.040bB	create a new genus in the family *Siphoviridae*, order *Caudovirales*
2015.040cB	name the genus *Slashvirus*
2015.040dB	designate *Bacillus virus Slash* as type species of the new genus

2015.041a-dB.A.v2.Wphvirus [41]

2015.041aB	create species in genus *Wphvirus* (new) in the family *Myoviridae*: *Bacillus virus WPh*
2015.041bB	create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.041cB	name the genus *Wphvirus*
2015.041dB	designate *Bacillus virus WPh* as type species of the new genus

2015.042a-oB.A.v3.Pleolipoviridae [42]

2015.042aB	create 5 species in genus *Alphapleolipovirus* (new), family *Pleolipoviridae* (new): *Halorubrum virus HRPV-1, Halorubrum virus HRPV-2, Halorubrum virus HRPV-6, Haloarcula virus HHPV-1, Haloarcula virus HHPV-2*
2015.042bB	create a new genus in the new family *Pleolipoviridae*
2015.042cB	name the genus *Alphapleolipovirus*
2015.042dB	designate *Halorubrum virus HRPV-1* as type species of the new genus
2015.042eB	create 2 species in genus *Betapleolipovirus* (new), family *Pleolipoviridae* (new): *Halorubrum virus HRPV-3, Halogeometricum virus HGPV-1*
2015.042fB	create a new genus in the new family *Pleolipoviridae*
2015.042gB	name the genus *Betapleolipovirus*
2015.042hB	designate *Halorubrum virus HRPV-3* as type species of the new genus
2015.042iB	create species in genus *Gammapleolipovirus* (new), family *Pleolipoviridae* (new): *Haloarcula virus His2*
2015.042jB	create a new genus in the new family *Pleolipoviridae*
2015.042kB	name the genus *Gammapleolipovirus*
2015.042lB	designate *Haloarcula virus His2* as type species of the new genus

2015.042mB   create a new unassigned family
2015.042nB   name the family *Pleolipoviridae*
2015.042oB   assign 3 new genera (*Alphapleovirus*, *Betapleovirus* and *Gammapleovirus*) to the new family

2015.043a-dB.A.v2.Ap22virus [43]

2015.043aB   create 4 species in genus *Ap22virus* (new) in the family *Myoviridae*: *Acinetobacter virus AbC62, Acinetobacter virus AB2, Acinetobacter virus AB1, Acinetobacter virus AP22*
2015.043bB   create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.043cB   name the genus *Ap22virus*
2015.043dB   designate *Acinetobacter virus AP22* as type species of the new genus

2015.044a-nB.A.v1.Guernseyvirinae [44]

2015.044aB   create 6 species in genus *Jerseylikevirus* (proposed name: *Jerseyvirus*) in the family *Siphoviridae*: *Salmonella virus AG11, Salmonella virus SETP7, Salmonella virus SP101, Salmonella virus SETP13, Salmonella phage LSPA1, Salmonella phage L13*
2015.044bB   create species in the genus *Sp31virus* (new) subfamily *Guernseyvirinae* (new), family *Siphoviridae*: *Salmonella virus SP31*
2015.044cB   create a new genus in the subfamily *Guernseyvirinae* (new), family *Siphoviridae*, order *Caudovirales*
2015.044dB   name the genus *Sp31virus*
2015.044eB   designate *Salmonella virus SP31* as type species of the new genus
2015.044fB   create 4 species in the genus *K1gvirus* (new) subfamily *Guernseyvirinae* (new), family *Siphoviridae*: *Escherichia virus K1ind1, Escherichia virus K1G, Escherichia virus K1ind2, Escherichia virus K1H*
2015.044gB   create a new genus in the subfamily *Guernseyvirinae* (new), family *Siphoviridae*, order *Caudovirales*
2015.044hB   name the genus *K1gvirus*
2015.044iB   designate *Escherichia virus K1G* as type species of the new genus
2015.044jB   create a new subfamily in the family *Siphoviridae*, order *Caudovirales*
2015.044kB   name the subfamily *Guernseyvirinae*
2015.044lB   assign 3 genera [*Jerseylikevirus* (proposed name *Jerseyvirus*), *Sp31virus* (new) and *K1gvirus* (new)] to the new subfamily

2015.044mB   remove genus *Jerseylikevirus* from its position of unassigned in the family *Siphoviridae*
2015.044nB   reassign the above genus to subfamily *Guernseyvirinae* (new), family *Siphoviridae*

2015.045a-dBA.v2.Rheph4virus [45]

2015.045aB   create species *Rhizobium virus RHEph4* in genus *Rheph4virus* (new) in the family *Myoviridae*: *Rhizobium virus RHEph4*
2015.045bB   create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.045cB   name the genus *Rheph4virus*
2015.045dB   designate *Rhizobium virus RHEph4* as type species of the new genus

2015.047aB.A.v2.Felixo1virus_7sp [46]

2015.047aB   create 7 species in genus *Felixounalikevirus* (proposed name *Felixo1virus*), family *Myoviridae*, order *Caudovirales*: *Salmonella virus UAB87, Escherichia virus VpaE1, Salmonella virus Mushroom, Escherichia virus JH2, Escherichia virus AYO145A, Salmonella virus HB2014, Escherichia virus EC6*

2015.048a-dB.A.v2.Ssp2virus [47]

2015.048aB   create 2 species in the genus *Ssp2virus* (new), family *Siphoviridae*: *Vibrio virus SSP002, Vibrio virus MAR10*
2015.048bB   create a new genus in the family *Siphoviridae*, order *Caudovirales*
2015.048cB   name the genus *Ssp2virus*
2015.048dB   designate *Vibrio virus SSP002* as type species of the new genus

2015.049a-dB.A.v2.Cvm10virus [48]

2015.049aB   create 2 species in genus *Cvm10virus* (new) in the family *Myoviridae*: *Escherichia virus CVM10, Escherichia virus ep3*
2015.049bB   create a new genus in the family *Myoviridae*, order *Caudovirales*
2015.049cB   name the genus *Cvm10virus*
2015.049dB   designate *Escherichia virus CVM10* as type species of the new genus

2015.050a-dB.A.v2.G7cvirus [49]

2015.050aB   create 8 species in genus *G7cvirus* (new) in the family *Podoviridae*: *Escherichia virus G7C, Escherichia virus APEC7, Escherichia virus APEC5, Escherichia virus Bp4, Escherichia virus ECBP1, Escherichia*

virus IME11, *Escherichia virus EC1UPM*,
Shigella virus Sb1

2015.050bB	create a new genus in the family *Podoviridae*, order *Caudovirales*
2015.050cB	name the genus *G7cvirus*
2015.050dB	designate *Escherichia virus G7C* as type species of the new genus

2015.051a-dB.A.v2.Nonagvirus [50]

2015.051aB	create 4 species in genus *Nonagvirus* (new) in the family *Siphoviridae*: *Escherichia virus 9g, Escherichia virus JenK1, Escherichia virus JenP2, Escherichia virus JenP1*
2015.051bB	create a new genus in the family *Siphoviridae*, order *Caudovirales*
2015.051cB	name the genus *Nonagvirus*
2015.051dB	designate *Escherichia virus 9g* as type species of the new genus

2015.052aB.A.v3.Schizot4virus_sp [51]

2015.052aB	create species in genus *Schizot4likevirus* (proposed name *Schizot4virus*), subfamily *Tevenvirinae*, family *Myoviridae*, order *Caudovirales*: *Vibrio virus ValKK3*

2015.053a-dB.A.v2.Seuratvirus [52]

2015.053aB	create 2 species in genus *Seuratvirus* (new) in the family *Siphoviridae*: *Escherichia virus Seurat, Escherichia virus CAjan*
2015.053bB	create a new genus in the family *Siphoviridae*, order *Caudovirales*
2015.053cB	name the genus *Seuratvirus*
2015.053dB	designate *Escherichia virus Seurat* as type species of the new genus

2015.054a-dB.A.v3.Septima3virus [53]

2015.054aB	create 4 species in genus *Septima3virus* (new) in the family *Siphoviridae*: *Pseudomonas virus 73, Pseudomonas virus Kakheti25, Pseudomonas virus Ab26, Burkholderia virus KL1*
2015.054bB	create a new genus in the family *Siphoviridae*, order *Caudovirales*
2015.054cB	name the genus *Septima3virus*
2015.054dB	designate *Pseudomonas virus 73* as type species of the new genus

Animal DNA Viruses and Retroviruses (code D or V)
2015.001aD.A.v2.Betabaculovirus_sp [54]

2015.001aD	create species in the genus *Betabaculovirus*, family *Baculoviridae*: *Agrotis segetum granulovirus*

2015.002a,bD.A.v2.Alphabaculovirus_sp,ren [55]

2015.002aD	create species in genus *Alphabaculovirus*, family *Baculoviridae*: *Agrotis segetum nucleopolyhedrovirus B*
2015.002bD	in genus *Alphabaculovirus* change the name of species *Agrotis segetum nucleopolyhedrovirus* to *Agrotis segetum nucleopolyhedrovirus A*

2015.003a,bD.A.v1.Kappatorquevirus_sp,ren [56]

2015.003aD	create species in genus *Kappatorquevirus* (family *Anelloviridae*): *Torque teno sus virus k2b*
2015.003bD	in genus *Kappatorquevirus* (family *Anelloviridae*), change the name of species *Torque teno sus virus k2* to *Torque teno sus virus k2a*

2014.006a,hD.A.v6.Circovirus_11spren [57]

2014.006aV	create 11 species in the genus *Circovirus*, family *Circoviridae*: *Barbel circovirus, Bat circovirus 1, Bat circovirus 2, Bat circovirus 3, Canine circovirus, Chimpanzee faeces associated circovirus, European catfish circovirus, Human faeces associated circovirus, Mink circovirus, Raven circovirus, Zebra finch circovirus*
2014.006hV	change the name of species *Porcine circovirus-1* to *Porcine circovirus 1* and *Porcine circovirus-2* to *Porcine circovirus 2*

2014.006b-eD.A.v3.Cyclovirus [58]

2014.006bV	create 28 species in the genus *Cyclovirus* (new), family *Circoviridae*: *Bat cyclovirus, Bat faeces associated cyclovirus 1, Bat faeces associated cyclovirus 2, Bat faeces associated cyclovirus 3, Bat faeces associated cyclovirus 4, Bovine cyclovirus, Chimpanzee faeces associated cyclovirus, Dragonfly cyclovirus 1, Dragonfly cyclovirus 2, Dragonfly cyclovirus 3, Dragonfly cyclovirus 4, Dragonfly cyclovirus 5, Dragonfly cyclovirus 6, Dragonfly cyclovirus 7, Dragonfly cyclovirus 8, Florida wood cockroach cyclovirus, Gallus cyclovirus, Goat cyclovirus, Human cyclovirus 1, Human cyclovirus 2, Human cyclovirus 3, Human faeces associated cyclovirus 1, Human*

faeces associated cyclovirus 2, Human faeces associated cyclovirus 3, Human faeces associated cyclovirus 4, Human faeces associated cyclovirus 5, Human faeces associated cyclovirus 6, Human faeces associated cyclovirus 8

2014.006cV	create a new genus in the family *Circoviridae*
2014.006dV	name the genus *Cyclovirus*
2014.006eV	designate *Human cyclovirus 1* as type species of the new genus

2014.006f,gD.A.v3.Gyrovirus_move [59]

2014.006fV	remove the genus *Gyrovirus* and constituent species from family *Circoviridae*
2014.006gV	reassign genus *Gyrovirus* to the familly *Anelloviridae*

2015.004a-eD.A.v3.Toursvirus [60]

2015.004aD	create a new genus in family *Ascoviridae*
2015.004bD	name the genus *Toursvirus*
2015.004cD	designate *Diadromus pulchellus ascovirus 4a* as type species of the new genus
2015.004dD	remove species *Diadromus pulchellus ascovirus 4a* from genus *Ascovirus*
2015.004eD	re-assign the above species to genus *Toursvirus*

2015.005aD.A.v2.Aviadenovirus_4sp [61]

2015.005aD	create 4 species in the genus *Aviadenovirus*, family *Adenoviridae*: *Duck aviadenovirus B, Pigeon aviadenovirus A, Turkey aviadenovirus C, Turkey aviadenovirus D*

2015.006aD.A.v2.Simplexvirus_sp [62]

2015.006aD	create species in genus *Simplexvirus*, subfamily *Alphaherpesvirinae*, family *Herpesviridae*: *Panine alphaherpesvirus 3*

2015.007aD.A.v2.Alphabaculovirus_sp [63]

2015.007aD	create species in genus *Alphabaculovirus*, family *Baculoviridae*: *Chrysodeixis includens nucleopolyhedrovirus*

2015.009a,bD.A.v3.Orthohepadnavirus_2sp [64]

2015.009aD	create 2 species in the genus *Orthohepadnavirus*, family *Hepadnaviridae*: *Tent-making bat hepatitis B virus, Roundleaf bat hepatitis B virus*
2015.009bD	change the name of species *Bat hepatitis virus* (genus *Orthohepadnavirus*) to *Long-fingered bat hepatitis B virus*

2015.010aD.A.v2.Herpesviridae_spren [65]

2015.010aD	change species names in the family *Herpesviridae* to a binomial format based on the name of the subfamily (example: species *Gallid herpesvirus 1* [genus *ltovirus*, subfamily *Alphaherpesvirinae*] changed to *Gallid alphaherpesvirus 1*). For a full list of proposed changes, please refer to the full proposal

2015.011aD.A.v2.Betabaculovirus_sp [66]

2015.011aD	create species in genus *Betabaculovirus*, family *Baculoviridae*: *Erinnyis ello granulovirus*

2015.012aD.A.v2.Mastadenovirus_2sp [67]

2015.012aD	create 2 species in the genus *Mastadenovirus*, family *Adenoviridae*: *Simian mastadenovirus B, Simian mastadenovirus C*

2015.013aD.A.v2.Betabaculovirus_sp [68]

2015.013aD	create species in genus *Betabaculovirus*, family *Baculoviridae*: *Spodoptera frugiperda granulovirus*

2015.014a-atD.A.v2.Papillomaviridae_10gen [69]

2015.014aD	create 6 species in genus *Gammapapillomavirus*, family *Papillomaviridae*: *Gammapapillomavirus 21, Gammapapillomavirus 22, Gammapapillomavirus 23, Gammapapillomavirus 24, Gammapapillomavirus 25, Gammapapillomavirus 26*
2015.014bD	create species in genus *Mupapillomavirus*, family *Papillomaviridae*: *Mupapillomavirus 3*
2015.014cD	create species in genus *Xipapillomavirus*, family *Papillomaviridae*: *Xipapillomavirus 3*
2015.014dD	create species in genus *Rhopapillomavirus*, family *Papillomaviridae*: *Rhopapillomavirus 2*
2015.014eD	create species in genus *Taupapillomavirus*, family *Papillomaviridae*: *Taupapillomavirus 3*
2015.014fD	create species in genus *Dyokappapapillomavirus*, family *Papillomaviridae*: *Dyokappapapillomavirus 2*

2015.014gD	create species in genus *Dyotaupapillomavirus* (new), family *Papillomaviridae*: *Dyotaupapillomavirus 1*
2015.014hD	create species in genus *Dyoupsilonpapillomavirus* (new), family *Papillomaviridae*: *Dyoupsilonpapillomavirus 1*
2015.014iD	create species in genus *Dyophipapillomavirus* (new), family *Papillomaviridae*: *Dyophipapillomavirus 1*
2015.014jD	create species in genus *Dyochipapillomavirus* (new), family *Papillomaviridae*: *Dyochipapillomavirus 1*
2015.014kD	create species in genus *Dyopsipapillomavirus* (new), family *Papillomaviridae*: *Dyopsipapillomavirus 1*
2015.014lD	create species in genus *Dyoomegapapillomavirus* (new), family *Papillomaviridae*: *Dyoomegapapillomavirus 1*
2015.014mD	create species in genus *Treisdeltapapillomavirus* (new), family *Papillomaviridae*: *Treisdeltapapillomavirus 1*
2015.014nD	create species in genus *Treisepsilonpapillomavirus* (new), family *Papillomaviridae*: *Treisepsilonpapillomavirus 1*
2015.014oD	create species in genus *Treiszetapapillomavirus* (new), family *Papillomaviridae*: *Treiszetapapillomavirus 1*
2015.014pD	create species in genus *Treisetapapillomavirus* (new), family *Papillomaviridae*: *Treisetapapillomavirus 1*
2015.014qD	create new genus in the family *Papillomaviridae*
2015.014rD	name the genus *Taupapillomavirus*
2015.014sD	designate *Taupapillomavirus 1* as the type species of the new genus
2015.014tD	create new genus in the family *Papillomaviridae*
2015.014uD	name the genus *Dyoupsilonpapillomavirus*
2015.014vD	designate *Dyoupsilonpapillomavirus 1* as the type species of the new genus
2015.014wD	create new genus in the family *Papillomaviridae*

2015.014xD	name the genus *Dyophipapillomavirus*
2015.014yD	designate *Dyophipapillomavirus 1* as the type species of the new genus
2015.014zD	create new genus in the family *Papillomaviridae*
2015.014aaD	name the genus *Dyochipapillomavirus*
2015.014abD	designate *Dyochipapillomavirus 1* as the type species of the new genus
2015.014acD	create new genus in the family *Papillomaviridae*
2015.014adD	name the genus *Dyopsipapillomavirus*
2015.014aeD	designate *Dyopsipapillomavirus 1* as the type species of the new genus
2015.014afD	create new genus in the family *Papillomaviridae*
2015.014agD	name the genus *Dyoomegapapillomavirus*
2015.014ahD	designate *Dyoomegapapillomavirus 1* as the type species of the new genus
2015.014aiD	create new genus in the family *Papillomaviridae*
2015.014ajD	name the genus *Treisdeltapapillomavirus*
2015.014akD	designate *Treisdeltapapillomavirus 1* as the type species of the new genus
2015.014alD	create new genus in the family *Papillomaviridae*
2015.014amD	name the genus *Treisepsilonpapillomavirus*
2015.014anD	designate *Treisepsilonpapillomavirus 1* as the type species of the new genus
2015.014aoD	create new genus in the family *Papillomaviridae*
2015.014apD	name the genus *Treiszetapapillomavirus*
2015.014aqD	designate *Treiszetapapillomavirus 1* as the type species of the new genus
2015.014arD	create new genus in the family *Papillomaviridae*
2015.014asD	name the genus *Treisetapapillomavirus*
2015.014atD	designate *Treisetapapillomavirus 1* as the type species of the new genus

2015.015a-aaD.A.v2.Polyomaviridae_rev [70]

2015.015aD	create 34 species in genus *Alphapolyomavirus* (new), family *Polyomaviridae*: *Acerodon celebensis polyomavirus 1, Artibeus planirostris polyomavirus 2, Artibeus planirostris polyomavirus 3, Ateles paniscus polyomavirus 1, Cardioderma cor polyomavirus 1, Carollia perspicillata polyomavirus 1, Chlorocebus pygerythrus polyomavirus 1, Chlorocebus pygerythrus polyomavirus 3, Dobsonia moluccensis polyomavirus 1, Eidolon helvum*

polyomavirus 1, Gorilla gorilla polyomavirus 1, Human polyomavirus 5, Human polyomavirus 8, Human polyomavirus 9, Human polyomavirus 12, Human polyomavirus 13, Macaca fascicularis polyomavirus 1, Molossus molossus polyomavirus 1, Otomops martiensseni polyomavirus 1, Otomops martiensseni polyomavirus 2, Pan troglodytes polyomavirus 1, Pan troglodytes polyomavirus 2, Pan troglodytes polyomavirus 3, Pan troglodytes polyomavirus 4, Pan troglodytes polyomavirus 5, Pan troglodytes polyomavirus 6, Pan troglodytes polyomavirus 7, Papio cynocephalus polyomavirus 1, Piliocolobus rufomitratus polyomavirus 1, Pongo abelii polyomavirus 1, Pongo pygmaeus polyomavirus 1, Procyon lotor polyomavirus 1, Pteropus vampyrus polyomavirus 1, Sturnira lilium polyomavirus 1

2015.015bD create 22 species in genus *Betapolyomavirus* (new), family *Polyomaviridae: Acerodon celebensis polyomavirus 2, Artibeus planirostris polyomavirus 1, Cebus albifrons polyomavirus 1, Cercopithecus erythrotis polyomavirus 1, Chlorocebus pygerythrus polyomavirus 2, Desmodus rotundus polyomavirus 1, Dobsonia moluccensis polyomavirus 2, Dobsonia moluccensis polyomavirus 3, Equus caballus polyomavirus 1, Human polyomavirus 3, Human polyomavirus 4, Loxodonta africana polyomavirus 1, Mastomys natalensis polyomavirus 1, Meles meles polyomavirus 1, Miniopterus africanus polyomavirus 1, Myotis lucifugus polyomavirus 1, Papio cynocephalus polyomavirus 2, Pteronotus davyi polyomavirus 1, Pteronotus parnellii polyomavirus 1, Saimiri boliviensis polyomavirus 1, Saimiri sciureus polyomavirus 1, Zalophus californianus polyomavirus 1*

2015.015cD create 6 species in genus *Gammapolyomavirus* (new), family *Polyomaviridae: Anser anser polyomavirus 1, Corvus monedula polyomavirus 1, Cracticus torquatus*

polyomavirus 1, Pygoscelis adeliae polyomavirus 1, Pyrrhula pyrrhula polyomavirus 1, Serinus canaria polyomavirus 1

2015.015dD create 4 species in genus *Deltapolyomavirus* (new), family *Polyomaviridae: Human polyomavirus 6, Human polyomavirus 7, Human polyomavirus 10, Human polyomavirus 11*

2015.015eD create new genus in the family *Polyomaviridae*

2015.015fD name the genus *Alphapolyomavirus*

2015.015gD designate *Mus musculus polyomavirus 1* as the type species of the new genus

2015.015hD create new genus in the family *Polyomaviridae*

2015.015iD name the genus *Betapolyomavirus*

2015.015jD designate *Macaca mulatta polyomavirus 1* as the type species of the new genus

2015.015kD create new genus in the family *Polyomaviridae*

2015.015lD name the genus *Gammapolyomavirus*

2015.015mD designate *Aves polyomavirus 1* as the type species of the new genus

2015.015nD create new genus in the family *Polyomaviridae*

2015.015oD name the genus *Deltapolyomavirus*

2015.015pD designate *Human polyomavirus 6* as the type species of the new genus

2015.015qqD remove the genus *Polyomavirus*

2015.015rD remove 4 species (*African green monkey polyomavirus, Simian virus 12, Baboon polyomavirus 2, Rabbit kidney vacuolating virus*) from their unassigned position in the family *Polyomaviridae*

2015.015sD remove species *Murine polyomavirus* and *Hamster polyomavirus* from genus *Polyomavirus* (to be removed) in the family *Polyomaviridae*

2015.015tD reassign the above species to genus *Alphapolyomavirus* (new), family *Polyomaviridae*

2015.015uD remove 4 species (*BK polyomavirus, JC polyomavirus, Murine pneumotropic virus, Simian virus 40*) from genus *Polyomavirus* (to be removed) in the family *Polyomaviridae*

2015.015vD reassign the above species to genus *Betapolyomavirus* (new), family *Polyomaviridae*

2015.015wD remove species *Budgerigar fledgling disease polyomavirus* from genus

Polyomavirus (to be removed) in the family *Polyomaviridae*

2015.015xD   reassign the above species to genus *Gammapolyomavirus* (new), family *Polyomaviridae*

2015.015yD   remove species *Bovine polyomavirus* from genus *Polyomavirus* (to be removed) in the family *Polyomaviridae*

2015.015zD   reassign the above species to genus *Deltapolyomavirus* (new), family *Polyomaviridae*

2015.015aaD   change the names of the following 8 species: *BK polyomavirus* to *Human polyomavirus 1*; *Bovine polyomavirus* to *Bos taurus polyomavirus 1*; *Budgerigar fledgling disease polyomavirus* to *Aves polyomavirus 1*; *Hamster polyomavirus* to *Mesocricetus auratus polyomavirus 1*; *JC polyomavirus* to *Human polyomavirus 2*; *Murine polyomavirus* to *Mus musculus polyomavirus 1*; *Murine pneumotropic virus* to *Mus musculus polyomavirus 2*; *Simian virus 40* to *Macaca mulatta polyomavirus 1*

2015.016aD.A.v2.Alphabaculovirus_sp [71]

2015.016aD   create species in the genus *Alphabaculovirus*, family *Baculoviridae*: *Sucra jujuba nucleopolyhedrovirus*

2015.001a-kF.A.v1.Lavidaviridae [72]

2015.001aF   create 2 species in genus *Sputnikvirus* (new), family *Lavidaviridae* (new): *Mimivirus-dependent virus Sputnik, Mimivirus-dependent virus Zamilon*

2015.001bF   create species in genus *Mavirus* (new), family *Lavidaviridae* (new): *Cafeteriavirus-dependent mavirus*

2015.001cF   create a new genus in the family *Lavidaviridae* (new)

2015.001dF   name the new genus *Sputnikvirus*

2015.001eF   designate *Mimivirus-dependent virus Sputnik* as type species in the new genus

2015.001fF   create a new genus in the family *Lavidaviridae* (new)

2015.001gF   name the new genus *Mavirus*

2015.001hF   designate *Cafeteriavirus-dependent mavirus* as type species in the new genus

2015.001iF   create a new unassigned family

2015.001jF   name the new family *Lavidaviridae*

2015.001kF   assign 2 genera, *Sputnikvirus* (new) and *Mavirus* (new), to the new family

2015.001dF   name the new genus *Sputnikvirus*
2015.001eF
2015.001fF
2015.001gF
2015.001hF
2015.001iF
2015.001jF

2015.001kF   designate *Mimivirus-dependent virus Sputnik* as type species in the new genus create a new genus in the family *Lavidaviridae* (new) name the new genus *Mavirus* designate *Cafeteriavirus-dependent mavirus* as type species in the new genus create a new unassigned family name the new family *Lavidaviridae* assign 2 genera, *Sputnikvirus* (new) and *Mavirus* (new), to the new family

Fungal and Protist Viruses (code F)
2014.001a-dF.A.v3.Botybirnaviridae [73]

2014.001aF   create species in the unassigned genus *Botybirnavirus* (new): *Botrytis porri botybirnavirus 1*

2014.001bF   create new unassigned genus

2014.001cF   name the genus *Botybirnavirus*

2014.001dF   designate *Botrytis porri botybirnavirus 1* as the type species of the new genus

2015.002a-gF.A.v2.Genomoviridae [74]

2015.002aF   create species in genus *Gemycircularvirus* (new), family *Genomoviridae* (new): *Sclerotinia gemycircularvirus 1*

2015.002bF   create a new genus in the family *Genomoviridae* (new)

2015.002cF   name the new genus *Gemycircularvirus*

2015.002dF   designate *Sclerotinia gemycircularvirus 1* as type species in the new genus

2015.002eF   create a new unassigned family

2015.002fF   name the new family *Genomoviridae*

2015.002gF   assign genus *Gemycircularvirus* (new), to the new family

2015.003a,bF.A.v3.Partitiviridae_4sp [75]

2015.003aF   create 3 species in genus *Alphapartitivirus*, family *Partitiviridae*: *Heterobasidion partitivirus 12, Heterobasidion partitivirus 13, Heterobasidion partitivirus 15*

2015.003bF   create species in genus *Betapartitivirus*, family *Partitiviridae*: *Heterobasidion partitivirus 7*

General Issues (code G)
2015.001aG.A.v1.Diacritics [76]

2015.001aG   Modify the ICVCN, Section II, paragraph 3.13, to disallow diacritics and other non-standard characters in taxon names

2015.002aG.A.v2.Abolish_Rule_3.35 [77]

2015.002aG   Abolish ICVCN Rule 3.35 to allow classification of satellite viruses and other virus-related agents

Animal dsRNA and ssRNA- Viruses (code M or V)
2014.003a-dV.A.v3.Dichorhavirus [78]

2014.003aV   create 2 species in the genus *Dichorhavirus* (new), family *Rhabdoviridae*, order *Mononegavirales*: *Orchid fleck dichorhavirus, Coffee ringspot dichorhavirus*

2014.003bV   create a new genus in the family *Rhabdoviridae*, order *Mononegavirales*

2014.003cV   name the genus *Dichorhavirus*

2014.003dV   designate *Orchid fleck dichorhavirus* as type species of the new genus

2015.001aM.A.v2.Mammarenavirus_4sp [79]

2015.001aM   create 4 species in the genus *Mammarenavirus*, family *Arenaviridae*: *Gairo mammarenavirus, Mariental mammarenavirus, Okahandja mammarenavirus, Wenzhou mammarenavirus*

2015.002aM.A.v2.Bornavirus_2sp [80]

2015.002aM   create 2 species in the genus *Bornavirus*, family *Bornaviridae*, order *Mononegavirales*: *Elapid 1 bornavirus, Psittaciform 2 bornavirus*

2015.003aM.A.v5.Bunyaviridae_spren [81]

2015.003aM   convert all species names in the family *Bunyaviridae* to a non-Latinized binomial format [example: *Andes virus* (in the genus *Hantavirus*) changed to *Andes hantavirus*]. For a complete list of existing and proposed names, please refer to the original proposal

2015.004a-fM.A.v3.Socyvirus [82]

2015.004aM   create a new genus in the family *Nyamiviridae*, order *Mononegavirales*

2015.004bM   name the new genus *Socyvirus*

2015.004cM   designate *Soybean cyst nematode socyvirus* (new name) as type species in the new genus

2015.004dM   remove species *Soybean cyst nematode socyvirus* (new name) from its unassigned position in the family *Nyamiviridae*

2015.004eM   reassign the above species to genus *Socyvirus* (new)

2015.004fM   change the name of the unassigned species *Soybean cyst nematode virus* in the family *Nyamiviridae* to *Soybean cyst nematode socyvirus*

2015.005a,bM.A.v1.Varicosavirus_ass [83]

2015.005aM   remove genus *Varicosavirus* from its unassigned (floating) position

2015.005bM   reassign the above genus to the family *Rhabdoviridae*, order *Mononegavirales*

2015.006aM.A.v5.Rhabdoviridae_spren [84]

2015.006aM   convert all species names in the family *Rhabdoviridae* to a non-Latinized binomial format [example: *Barley yellow striate mosaic virus* (in the genus *Cytorhabdovirus*) changed to *Barley yellow striate mosaic cytorhabdovirus*]. For a complete list of existing and proposed names, please refer to the original proposal

2015.007aM.A.v1.Cytorhabdovirus_sp [85]

2015.007aM   create species in the genus *Cytorhabdovirus*, family *Rhabdoviridae*: *Alfalfa dwarf cytorhabdovirus*

2015.008a-gM.A.v3.Mymonaviridae [86]

2015.008aM   create species in genus *Sclerotimonavirus* (new), family *Mymonaviridae* (new), order *Mononegavirales*: *Sclerotinia sclerotimonavirus*

2015.008bM   create a new genus in the family *Mymonaviridae* (new)

2015.008cM   name the new genus *Sclerotimonavirus*

2015.008dM   designate *Sclerotinia sclerotimonavirus* as type species in the new genus

2015.008eM   create a new family in the order *Mononegavirales*

2015.008fM   name the new family *Mymonaviridae*

2015.008gM   assign genus *Sclerotimonavirus* (new) to the new family

2015.009a-gM.A.v4.Sunviridae [87]

2015.009aM   create species in genus *Sunshinevirus* (new), family *Sunviridae* (new), order *Mononegavirales*: *Reptile sunshinevirus 1*

2015.009bM   create a new genus in the family *Sunviridae* (new) order *Mononegavirales*

2015.009cM	name the new genus *Sunshinevirus*
2015.009dM	designate *Reptile sunshinevirus 1* as type species in the new genus
2015.009eM	create a new family in the order *Mononegavirales*
2015.009fM	name the new family *Sunviridae*
2015.009gM	assign genus *Sunshinevirus* (new) to the new family

2015.010a-tM.A.v3.Mononegavirales_5gen [88]

2015.010aM	create species in genus *Crustavirus* (new) unassigned in the order *Mononegavirales*: *Wenzhou crustavirus*
2015.010bM	create a new unassigned genus in the order *Mononegavirales*
2015.010cM	name the new genus *Crustavirus*
2015.010dM	designate *Wenzhou crustavirus* as type species in the new genus
2015.010eM	create species in genus *Anphevirus* (new) unassigned in the order *Mononegavirales*: *Xincheng anphevirus*
2015.010fM	create a new genus unassigned in the order *Mononegavirales*
2015.010gM	name the new genus *Anphevirus*
2015.010hM	designate *Xincheng anphevirus* as type species in the new genus
2015.010iM	create species in genus *Arlivirus* (new) unassigned in the order *Mononegavirales*: *Lishi arlivirus*
2015.010jM	create a new genus unassigned in the order *Mononegavirales*
2015.010kM	name the new genus *Arlivirus*
2015.010lM	designate *Lishi arlivirus* as type species in the new genus
2015.010mM	create species in genus *Wastrivirus* (new) unassigned in the order *Mononegavirales*: *Sanxia wastrivirus*
2015.010nM	create a new genus unassigned in the order *Mononegavirales*
2015.010oM	name the new genus *Wastrivirus*
2015.010pM	designate *Sanxia wastrivirus* as type species in the new genus
2015.010qM	create species in genus *Chengtivirus* (new) unassigned in the order *Mononegavirales*: *Tacheng chengtivirus*
2015.010rM	create a new genus unassigned in the order *Mononegavirales*
2015.010sM	name the new genus *Chengtivirus*
2015.010tM	designate *Tacheng chengtivirus* as type species in the new genus

2015.011a-gM.A.v2.Pneumoviridae [89]

2015.011aM	remove genera *Orthopneumovirus* (new name) and *Metapneumovirus* from subfamily *Pneumovirinae*, family *Paramyxoviridae*, order *Mononegavirales*
2015.011bM	reassign the above genera to the new family *Pneumoviridae*
2015.011cM	remove all 7 genera (*Aquaparamyxovirus*, *Avulavirus*, *Ferlavirus*, *Henipavirus*, *Morbillivirus*, *Respirovirus* and *Rubulavirus*) from the subfamily *Paramyxovirinae*
2015.011dM	reassign the above genera to the family *Paramyxoviridae*
2015.011eM	remove (abolish) subfamilies *Pneumovirinae* and *Paramyxovirinae* from family *Paramyxoviridae*, order *Mononegavirales*
2015.011fM	create a new family in the order *Mononegavirales*
2015.011gM	name the new family *Pneumoviridae*
2015.011hM	assign 2 existing genera [*Orthopneumovirus* (new name) and *Metapneumovirus*] to the new family
2015.011iM	change the name of genus *Pneumovirus* to *Orthopneumovirus*

2015.012aM.A.v3.Avulavirus_3sp [90]

| 2015.012aM | create 3 species in the genus *Avulavirus*, subfamily *Paramyxovirinae* [subfamily to be abolished (2015.011a-gM)], family *Paramyxoviridae*, order *Mononegavirales*: *Avian paramyxovirus 10*, *Avian paramyxovirus 11*, *Avian paramyxovirus 12* |

2015.013aM.A.v3.Henipavirus_3sp [91]

| 2015.013aM | create 3 species in the genus *Henipavirus*, family *Paramyxoviridae*, order *Mononegavirales*: *Cedar henipavirus*, *Ghanaian bat henipavirus*, *Mojiang henipavirus* |

2015.014aM.A.v2.Morbillivirus_sp [92]

| 2015.014aM | create species in genus *Morbillivirus*, family *Paramyxoviridae*, order *Mononegavirales*: *Feline morbillivirus* |

2015.015aM.A.v1.Respirovirus_sprem [93]

| 2015.015aM | remove (abolish) species *Simian virus 10* from genus *Respirovirus*, |

subfamily *Paramyxovirinae*, family *Paramyxoviridae*

2015.017aM.A.v1.Respirovirus_sp [94]

2015.017aM   create species in the genus *Respirovirus*, subfamily *Paramyxovirinae*, family *Paramyxoviridae*, order *Mononegavirales*: *Porcine parainfluenza virus 1*

2015.018aM.A.v2.Fijivirus_sp [95]

2015.018aM   create species in genus *Fijivirus*, subfamily *Spinareovirinae*, family *Reoviridae*: *Southern rice black-streaked dwarf virus*

2015.019aM.A.v3.Orthoreovirus_sp [96]

2015.019aM   create species in genus *Orthoreovirus*, subfamily *Spinareovirinae*, family *Reoviridae*: *Piscine orthoreovirus*

Plant Viruses (code P)
2014.006b-fP.A.v3.Pelarspovirus [97]

2014.006bP   create a new genus
2014.006cP   name the genus *Pelarspovirus*
2014.006dP   designate *Pelargonium line pattern virus* as type species of the new genus
2014.006eP   remove 5 species (*Elderberry latent virus, Pelargonium chlorotic ring pattern virus, Pelargonium line pattern virus, Pelargonium ringspot virus and Rosa rugosa leaf distortion virus*) from their present, unassigned, position in the family *Tombusviridae*
2014.006fP   re-assign the above 5 species to the new genus created in 2014.006b-cP

2015.001a-dP.A.v2.Goravirus [98]

2015.001aP   create 2 species in the genus *Goravirus* (new), family *Virgaviridae*: *Gentian ovary ringspot virus, Drakaea virus A*
2015.001bP   create a new genus in the family *Virgaviridae*
2015.001cP   name the genus *Goravirus*
2015.001dP   designate *Gentian ovary ringspot virus* as type species of the new genus

2015.002aP.A.v2.Poacevirus [99]

2015.002aP   create species in the genus *Poacevirus*, family *Potyviridae*: *Caladenia virus A*

2015.003a-dP.A.v2.Potyviridae [100]

2015.003aP   create 5 species in the genus *Potyvirus*, family *Potyviridae*: *Catharanthus mosaic virus, Donkey orchid virus A, Sunflower mild mosaic virus, Tamarillo leaf malformation virus, Vanilla distortion mosaic virus*
2015.003bP   remove species *Artichoke latent virus* from genus *Potyvirus*
2015.003cP   reassign above species to genus *Macluravirus*
2015.003dP   remove (abolish) species *Ranunculus latent virus* from genus *Macluravirus*

2015.005aP.A.v3.Nepovirus_2sp [101]

2015.005aP   create 2 species in genus *Nepovirus*, subfamily *Comovirinae*, family *Secoviridae*, order *Picornavirales*: *Aeonium ringspot virus, Mulberry mosaic leaf roll associated virus*

2015.006aP.A.v2.Torradovirus_2sp [102]

2015.006aP   create 2 species in genus *Torradovirus*, family *Secoviridae*, order *Picornavirales*: *Carrot torradovirus 1, Motherwort yellow mottle virus*

2015.007a-rP.A.v1.split_Carmovirus [103]

2015.007aP   create a new genus in the family *Tombusviridae*
2015.007bP   name the genus *Alphacarmovirus*
2015.007cP   designate *Carnation mottle virus* as type species of the new genus
2015.007dP   create a new genus in the family *Tombusviridae*
2015.007eP   name the genus *Betacarmovirus*
2015.007fP   designate *Turnip crinkle virus* as type species of the new genus
2015.007gP   create a new genus in the family *Tombusviridae*
2015.007hP   name the genus *Gammacarmovirus*
2015.007iP   designate *Melon necrotic spot virus* as type species of the new genus
2015.007jP   remove 7 species (*Angelonia flower break virus, Calibrachoa mottle virus, Carnation mottle virus, Honeysuckle ringspot virus, Nootka lupine vein clearing virus, Pelargonium flower break virus and Saguaro cactus virus*) from the genus *Carmovirus*, family *Tombusviridae*
2015.007kP   reassign the above 7 species to the genus *Alphacarmovirus* (new), family *Tombusviridae*
2015.007lP   remove 4 species (*Cardamine chlorotic fleck virus, Hibiscus chlorotic ringspot*

virus, Japanese iris necrotic ring virus and *Turnip crinkle virus*) from the genus *Carmovirus,* family *Tombusviridae*

2015.007mP   reassign the above 4 species to the genus *Betacarmovirus* (new), family *Tombusviridae*

2015.007nP   remove 4 species (*Cowpea mottle virus, Melon necrotic spot virus, Soybean yellow mottle mosaic virus* and *Pea stem necrosis virus* from the genus *Carmovirus,* family *Tombusviridae*

2015.007oP   reassign the above species to the genus *Gammacarmovirus* (new), family *Tombusviridae*

2015.007pP   remove 4 species (*Ahlum waterborne virus, Bean mild mosaic virus, Cucumber soil-borne virus* and *Weddel waterborne virus* from the genus *Carmovirus,* family *Tombusviridae*

2015.007qP   reassign the above species as unassigned in the family *Tombusviridae*

2015.007rP   remove genus *Carmovirus* from the family *Tombusviridae*

2015.008aP.A.v1.Aureusvirus_sp [104]

2015.008aP   create species in genus *Aureusvirus*, family *Tombusviridae*: *Yam spherical virus*

2015.009a-pP.A.v3.Plant_sat_4unassgen [105]

2015.009aP   create 3 species in the unassigned genus *Albetovirus* (new): *Tobacco albetovirus 1, Tobacco albetovirus 2, Tobacco albetovirus 3*

2015.009bP   create species in the unassigned genus *Aumaivirus* (new): *Maize aumaivirus 1*

2015.009cP   create species in the unassigned genus *Papanivirus* (new): *Panicum papanivirus 1*

2015.009dP   create species in the unassigned genus *Virtovirus* (new): *Tobacco virtovirus 1*

2015.009eP   create a new unassigned genus

2015.009fP   name the genus *Albetovirus*

2015.009gP   designate *Tobacco albetovirus 1* as type species of the new genus

2015.009hP   create a new unassigned genus

2015.009iP   name the genus *Aumaivirus*

2015.009jP   designate *Maize aumaivirus 1* as type species of the new genus

2015.009kP   create a new unassigned genus

2015.009lP   name the genus *Papanivirus*

2015.009mP   designate *Panicum papanivirus 1* as type species of the new genus

2015.009nP   create a new unassigned genus

2015.009oP   name the genus *Virtovirus*

2015.009pP   designate *Tobacco virtovirus 1* as type species of the new genus

2015.010aP.A.v1.Nanovirus_2sp [106]

2015.010aP   create two species in the genus *Nanovirus*, family *Nanoviridae*: *Black medic leaf roll virus, Pea yellow stunt virus*

2015.011a-adP.A.v2.Betaflexiviridae_rev [107]

2015.011aP   create two species in the genus *Chordovirus* (new), subfamily *Trivirinae* (new), family *Betaflexiviridae*, order *Tymovirales: Carrot Ch virus 1, Carrot Ch virus 2*

2015.011bP   create two species in the genus *Prunevirus* (new), subfamily *Trivirinae* (new), family *Betaflexiviridae*: *Apricot vein clearing associated virus, Caucasus prunus virus*

2015.011cP   create species in the genus *Tepovirus*, subfamily *Trivirinae* (new), family *Betaflexiviridae*: *Prunus virus T*

2015.011dP   create two species in the genus *Carlavirus*, subfamily *Quinvirinae* (new), family *Betaflexiviridae*: *Gaillardia latent virus, Potato virus H*

2015.011eP   create two species in the genus *Robigovirus* (new), subfamily *Quinvirinae* (new), family *Betaflexiviridae*: *Cherry rusty mottle associated virus, Cherry twisted leaf associated virus*

2015.011fP   create a new genus in the subfamily *Trivirinae* (new), family *Betaflexiviridae*, order *Tymovirales*

2015.011gP   name the genus *Chordovirus*

2015.011hP   designate *Carrot Ch virus 1* as type species of the new genus

2015.011iP   create a new genus in the subfamily *Trivirinae* (new), family *Betaflexiviridae*, order *Tymovirales*

2015.011jP   name the genus *Divavirus*

2015.011kP   designate *Diuris virus A* as type species of the new genus

2015.011lP   create a new genus in the subfamily *Trivirinae* (new), family *Betaflexiviridae*, order *Tymovirales*

2015.011mP   name the genus *Prunevirus*

2015.011nP   designate *Apricot vein clearing associated virus* as type species of the new genus

2015.011oP   create a new genus in the subfamily *Quinvirinae* (new), family *Betaflexiviridae*, order *Tymovirales*

2015.011pP	name the genus *Robigovirus*
2015.011qP	designate *Cherry necrotic rusty mottle virus* as type species of the new genus
2015.011rP	create a new subfamily within the family *Betaflexiviridae*, order *Tymovirales*
2015.011sP	name the subfamily *Trivirinae*
2015.011tP	assign the following 8 genera to the new subfamily: *Capillovirus, Citrivirus, Tepovirus, Trichovirus, Vitivirus, Chordovirus* (new), *Divavirus* (new) and *Prunevirus* (new)
2015.011uP	create a new subfamily within the family *Betaflexiviridae*, order *Tymovirales*
2015.011vP	name the subfamily *Quinvirinae*
2015.011wP	assign the following 3 genera to the new subfamily: *Carlavirus, Foveavirus* and *Robigovirus* (new)
2015.011xP	remove 3 species (*Diuris virus A, Diuris virus B* and *Hardenbergia virus A*) from their present, unassigned, position in the family *Betaflexiviridae*
2015.011yP	reassign the above 3 species to genus *Divavirus* (new), subfamily *Trivirinae* (new), family *Betaflexiviridae*, order *Tymovirales*
2015.011zP	remove 3 species (*African oil palm ringspot virus, Cherry green ring mottle virus* and *Cherry necrotic rusty mottle virus*) from their present, unassigned, position in the family *Betaflexiviridae*
2015.011aaP	reassign the above 3 species to genus *Robigovirus* (new), subfamily *Quinvirinae* (new), family *Betaflexiviridae*, order *Tymovirales*
2015.011abP	remove 3 species (*Banana mild mosaic virus, Banana virus X* and *Sugarcane striate mosaic-associated virus*) from their present, unassigned, position in the family *Betaflexiviridae*
2015.011acP	reassign the above 3 species to be unassigned in the subfamily *Quinvirinae* (new), family *Betaflexiviridae*, order *Tymovirales*
2015.011adP	remove (abolish) 7 species (*Dandelion latent virus, Elderberry symptomless virus, Honeysuckle latent virus, Hydrangea latent virus, Lilac mottle virus, Mulberry latent virus* and *Muskmelon vein necrosis virus*) from the genus *Carlavirus*, family *Betaflexiviridae*

2015.012a-dP.A.v5.Platypuvirus [108]

2015.012aP	create species in the genus *Platypuvirus* (new), family *Alphaflexiviridae*, order *Tymovirales*: *Donkey orchid symptomless virus*
2015.012bP	create a new genus in the family *Alphaflexiviridae*, order *Tymovirales*
2015.012cP	name the genus *Platypuvirus*
2015.012dP	designate *Donkey orchid symptomless virus* as type species of the new genus

2015.013aP.A.v3.Potexvirus_sp [109]

2015.013aP	create species in the genus *Potexvirus*, family *Alphaflexiviridae*, order *Tymovirales*: *Yam virus X*

2015.014aP.A.v1.Potexvirus_remsp [110]

2015.014aP	abolish 3 species (*Commelina virus X, Daphne virus X* and *Plantago severe mottle virus*) from the genus *Potexvirus*, family *Alphaflexiviridae*, order *Tymovirales*

2015.015a,bP.A.v3.Begomovirus_sprem [111]

2015.015aP	in the family *Geminiviridae*, create 57 species in genus *Begomovirus*: *Abutilon golden mosaic virus, Capraria yellow spot virus, Cassava mosaic Madagascar virus, Catharanthus yellow mosaic virus, Chenopodium leaf curl virus, Chilli leaf curl India virus, Chilli leaf curl Kanpur virus, Chilli leaf curl Vellanad virus, Clerodendron yellow mosaic virus, Clerodendrum golden mosaic China virus, Clerodendrum golden mosaic Jiangsu virus, Corchorus yellow vein mosaic virus, Cotton chlorotic spot virus, Crassocephalum yellow vein virus, Emilia yellow vein virus, French bean leaf curl virus, Hedyotis uncinella yellow mosaic virus, Hemidesmus yellow mosaic virus, Jatropha leaf curl virus, Jatropha mosaic Nigeria virus, Jatropha mosaic virus, Jatropha yellow mosaic virus, Malvastrum leaf curl Philippines virus, Mesta yellow vein mosaic Bahraich virus, Pepper yellow leaf curl virus, Pouzolzia golden mosaic virus, Pouzolzia mosaic Guangdong virus, Premna leaf curl virus, Rhynchosia yellow mosaic India virus, Sauropus leaf curl virus, Sida ciliaris golden*

mosaic virus, Sida common mosaic virus, Sida golden mosaic Brazil virus, Sida golden mosaic Lara virus, Sida yellow leaf curl virus, Sidastrum golden leaf spot virus, Soybean chlorotic blotch virus, Spinach yellow vein virus, Sunn hemp leaf distortion virus, Sweet potato leaf curl Henan virus, Sweet potato leaf curl Sichuan virus 1, Sweet potato leaf curl Sichuan virus 2, Tobacco leaf curl Comoros virus, Tomato bright yellow mosaic virus, Tomato bright yellow mottle virus, Tomato golden leaf distortion virus, Tomato interveinal chlorosis virus, Tomato leaf curl Liwa virus, Tomato leaf curl New Delhi virus 2, Tomato leaf curl New Delhi virus 4, Tomato leaf curl Palampur virus, Tomato leaf curl Patna virus, Tomato leaf curl Rajasthan virus, Tomato leaf curl Sulawesi virus, Velvet bean severe mosaic virus, Vigna yellow mosaic virus

2015.015bP   in the family *Geminiviridae*, remove the following 22 species from genus *Begomovirus*: *Abutilon Brazil virus, Ageratum leaf curl Cameroon virus, Bhendi yellow vein India virus, Bitter gourd yellow vein virus, Eclipta yellow vein virus, Hollyhock leaf crumple virus, Honeysuckle yellow vein Kagoshima virus, Kenaf leaf curl virus, Macroptilium yellow net virus, Malvastrum leaf curl Guangdong virus, Malvastrum yellow leaf curl virus, Malvastrum yellow vein Changa Manga virus, Okra leaf curl Cameroon virus, Okra yellow mottle Iguala virus, Okra yellow vein mosaic virus, Sida golden mosaic Honduras virus, Sida golden mosaic Liguanea virus, Sida yellow vein Madurai virus, Soybean chlorotic spot virus, Soybean crinkle leaf virus, Tomato leaf curl Kumasi virus, Tomato mosaic leaf curl virus*

2015.016aP.A.v2.Mastrevirus_3sp [112]

2015.016aP   create 3 species in genus *Mastrevirus* of the family *Geminiviridae*: *Sugarcane white streak virus, Axonopus compressus streak virus, Switchgrass mosaic-associated virus*

2015.017aP.A.v2.Sobemovirus_5sp [113]

2015.017aP   create 5 species in the unassigned genus *Sobemovirus*: *Artemisia virus A, Cymbidium chlorotic mosaic virus, Papaya lethal*

yellowing virus, Rottboellia yellow mottle virus, Soybean yellow common mosaic virus

2015.018aP.A.v3.Emaravirus_sp [114]

2015.018aP   create species in the unassigned genus, *Emaravirus*: *High Plains wheat mosaic virus*

2015.019aP.A.v2.Endornavirus_spa [115]

2015.019aP   create species in the genus *Endornavirus*, family *Endornaviridae*: *Basella alba endornavirus 1*

2015.020aP.A.v1.Endornavirus_spb [116]

2015.020aP   create species in the genus *Endornavirus*, family *Endornaviridae*: *Persea americana endornavirus 1*

2015.021aP.A.v1.Endornavirus_spc [117]

2015.021aP   create species in the genus *Endornavirus*, family *Endornaviridae*: *Yerba mate endornavirus 1*

2015.022aP.A.v1.Endornavirus_spd [118]

2015.022aP   create species in the genus *Endornavirus*, family *Endornaviridae*: *Sclerotinia sclerotiorum endornavirus 1*

2015.023aP.A.v2.Ophiovirus_sp [119]

2015.023aP   create species in the genus *Ophiovirus*, family *Ophioviridae*: *Blueberry mosaic associated virus*

2015.024a-cP.A.v2.Closteroviridae_5sp [120]

2015.024aP   create species in genus *Ampelovirus,* family *Closteroviridae*: *Blackberry vein banding-associated virus*
2015.024bP   create 3 species in genus *Velarivirus,* family *Closteroviridae*: *Cordyline virus 2, Cordyline virus 3, Cordyline virus 4*
2015.024cP   create an unassigned species in the family *Closteroviridae*: *Blueberry virus A*

2015.025aP.A.v1.Badnavirus_sp [121]

2015.025aP   create species in genus *Badnavirus*, family *Caulimoviridae*: *Grapevine Roditis leaf discoloration-associated virus*

2015.026aP.A.v2.Badnavirus_4sp [122]

2015.026aP   create 4 species in genus *Badnavirus*, family *Caulimoviridae*: *Sugarcane bacilliform Guadeloupe A virus, Sugarcane bacilliform*

Guadeloupe D virus, Cacao swollen shoot
Togo A virus, Cacao swollen shoot CD virus

Animal ssRNA+ Viruses (code S)
2015.002a-gS.A.v3.Sarthroviridae [123]

2015.002aS	create species in genus *Macronovirus* (new), family *Sarthroviridae* (new): *Macrobrachium satellite virus 1*
2015.002bS	create new genus in the family *Sarthroviridae* (new)
2015.002cS	name the genus *Macronovirus*
2015.002dS	designate *Macrobrachium satellite virus 1* as the type species of the new genus
2015.002eS	create a new unassigned family
2015.002fS	name the family *Sarthroviridae*
2015.002gS	assign genus *Macronovirus* (new) to the new family

2015.003a-eS.A.v2.Coronaviridae_12sp [124]

2015.003aS	create 3 species in genus *Alphacoronavirus*, subfamily *Coronavirinae*, family *Coronaviridae*, order *Nidovirales*: *Bat coronavirus HKU10, Bat coronavirus CDPHE15, Mink coronavirus 1*
2015.003bS	create 2 species in genus *Betacoronavirus*, subfamily *Coronavirinae*, family *Coronaviridae*, order *Nidovirales*: *Middle East respiratory syndrome-related coronavirus, Hedgehog coronavirus 1*
2015.003cS	create 5 species in genus *Deltacoronavirus*, subfamily *Coronavirinae*, family *Coronaviridae*, order *Nidoviarales*: *Coronavirus HKU15, White-eye coronavirus HKU16, Night heron coronavirus HKU19, Wigeon coronavirus HKU20, Common moorhen coronavirus HKU21*
2015.003dS	create species in genus *Bafinivirus*, subfamily *Torovirinae*, family *Coronaviridae*, order *Nidovirales*: *Fathead minnow nidovirus 1*
2015.003eS	create an unassigned species in the subfamily *Torovirinae*, family *Coronaviridae*, order *Nidovirales*: *Ball python nidovirus 1*

2015.004a,bS.A.v2.Mesoniviridae_6sp [125]

2015.004aS	create 4 species in genus *Alphamesonivirus*, family *Mesoniviridae*, order *Nidovirales*: *Alphamesonivirus 2, Alphamesonivirus 3, Alphamesonivirus 4, Alphamesonivirus 5*

2015.004bS	create 2 unassigned species in the family *Mesoniviridae*, order *Nidovirales*: *Mesonivirus 1, Mesonivirus 2*

2015.005a-dS.A.v2.Sinaivirus [126]

2015.005aS	create 2 species in the unassigned genus *Sinaivirus* (new): *Lake Sinai virus 1, Lake Sinai virus 2*
2015.005bS	create new unassigned genus
2015.005cS	name the genus *Sinai*virus
2015.005dS	designate *Lake Sinai virus 2* as the type species of the new genus

2015.006a-eS.A.v3.Triatovirus [127]

2015.006aS	create new genus in the family *Dicistroviridae*, order *Picornavirales*
2015.006bS	name the genus *Triatovirus*
2015.006cS	designate *Triatoma virus* as the type species of the new genus
2015.006dS	remove 5 species (*Triatoma virus, Plautia stali intestine virus, Himetobi P virus, Black queen cell virus, Homalodisca coagulata virus 1*) from genus *Cripavirus*, family *Dicistroviridae*
2015.006eS	reassign the above species to the new genus, *Triatovirus*

2015.007a-dS.A.v4.Limnipivirus [128]

2015.007aS	create 3 species in genus *Limnipivirus* (new), family *Picornaviridae*, order *Picornavirales*: *Limnipivirus A, Limnipivirus B, Limnipivirus C*
2015.007bS	create new genus in the family *Picornaviridae*, order *Picornavirales*
2015.007cS	name the genus *Limnipivirus*
2015.007dS	designate *Limnipivirus A* as the type species of the new genus

2015.008a-dS.A.v4.Potamipivirus [129]

2015.008aS	create species in genus *Potamipivirus* (new), family *Picornaviridae*, order *Picornavirales*: *Potamipivirus A*
2015.008bS	create new genus in the family *Picornaviridae*, order *Picornavirales*
2015.008cS	name the genus *Potamipivirus*
2015.008dS	designate *Potamipivirus A* as the type species of the new genus

2015.009aS.A.v2.Iflavirus_sp [130]

2015.009aS	create species in genus *Iflavirus*, family *Iflaviridae*, order *Picornavirales*: *Antheraea pernyi iflavirus*

2015.010aS.A.v2.Iflavirus_sp [131]

2015.010aS   create species in genus *Iflavirus*, family *Iflaviridae*, order *Picornavirales*: *Dinocampus coccinellae paralysis virus*

2015.011aS.A.v2.Iflavirus_sp [132]

2015.011aS   create species in genus *Iflavirus*, family *Iflaviridae*, order *Picornavirales*: *Lymantria dispar iflavirus 1*

2015.012aS.A.v2.Iflavirus_sp [133]

2015.012aS   create species in genus *Iflavirus*, family *Iflaviridae*, order *Picornavirales*: *Spodoptera exigua iflavirus 1*

2015.013aS.A.v2.Iflavirus_sp [134]

2015.013aS   create species in genus *Iflavirus*, family *Iflaviridae*, order *Picornavirales*: *Spodoptera exigua iflavirus 2*

2015.014a-cS.A.v3.Arteriviridae_sprev [135]

2015.014aS   create 9 species in genus *Arterivirus,* family *Arteriviridae*, order *Nidovirales*: *Porcine reproductive and respiratory syndrome virus 2, Kibale red-tailed guenon virus 1, Kibale red colobus virus 1, Kibale red colobus virus 2, Mikumi yellow baboon virus 1, Simian hemorrhagic encephalitis virus, DeBrazza's monkey arterivirus, Forest pouched giant rat arterivirus, Pebjah virus*

2015.014bS   create an unassigned species in the family *Arteriviridae*, order *Nidovirales*: *Wobbly possum disease virus*

2015.014dS   change the name of species *Porcine reproductive and respiratory syndrome virus* to *Porcine reproductive and respiratory syndrome virus 1*

Conclusion

All of the taxonomic proposals listed above were unanimously approved by the 80 members who voted (a return rate of 54%). The general proposal 2015.001aG [76] to modify the taxonomic code to disallow diacritics and other non-standard characters in taxon names was rejected by some but nevertheless received majority approval (54 in favour, 19 against and 7 abstentions). The changes listed above are now part of the official ICTV taxonomy and represent changes to the taxonomy published in the Ninth

Report [136] and are in addition to changes approved in February 2012 [137], February 2013 [138], March 2014 [139], and February 2015 [140]. 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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Create species *Agrotis segetum granulovirus* in the genus *Betabaculovirus*, family *Baculoviridae*. http://www.ictvonline.org/proposals-15/2015.001aD.A.v2.Betabaculovirus_sp.pdf

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Whitfield AE, Kuhn JH, Easton AJ, Nowotny N, Rima B, Rubbenstroth D, Walsh J, Clawson AN, Freitas-Astúa J, Goodin MM, Kitajima EW, Wetzel T (2015) ICTV taxonomic proposal 2014.003a-dV.A.v3.Dichorhavirus. Create 2 species, *Orchid fleck dichorhavirus* and *Coffee ringspot dichorhavirus*, in a new genus, *Dichorhavirus,* in the family *Rhabdoviridae*. http://www.ictvonline.org/proposals-15/2014.003a-dV.A.v3.Dichorhavirus.pdf

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family *Tombusviridae* assign 5 previously unassigned species to a new genus, *Pelarspovirus*. http://www.ictvonline.org/proposals-15/2014.006b-fP.A.v3.Pelarspovirus.pdf

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