

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* (*Myohalovirus*), *Phijlunlikevirus* (*Phijl1virus*), *Psimunlikevirus* (*Psimunavirus*), *Punalikevirus* (*P1virus*),
 2015.004cB name the genus *Ff47virus* (*Sfi1unlikevirus* (*Sfi11virus*),
 2015.004dB designate *Mycobacterium virus Ff47* as type species of the new genus *Sfi21dtunlikevirus* (*Sfi21dt1virus*),
Skunlikevirus (*Sk1virus*), *Spounlikevirus* (*Spo1virus*), *Tp2unlikevirus* (*Tp21virus*),
Tunlikevirus (*T1virus*), *Viunlikevirus* (*Vi1virus*)
- 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 *Caudovirales*
 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. *Andromedaviridae* to *Andromedavirus*). Also make the following individual name changes (proposed name in brackets):
3alikevirus (*Triavirus*), *77likevirus* (*Biseptimavirus*), *Ahjdlikevirus* (*P68virus*), *Bppunlikevirus* (*Bpp1virus*), *Cjwunlikevirus* (*Cjw1virus*), *Cp8unlikevirus* (*Cp8virus*), *Felixounlikevirus* (*Felixo1virus*), *Halolikevirus* (*Liefievirus*), *Hapunlikevirus* (*Hapunavirus*), *Hpunlikevirus* (*Hp1virus*), *I3likevirus* (*Bxz1virus*), *Iebhlikevirus* (*Cecivirus*), *Pbiunlikevirus* (*Pbi1virus*), *Pbunlikevirus* (*Pbunavirus*), *Pgonelikevirus* (*Pg1virus*), *Phic3unlikevirus* (*Phic31virus*), *Phicd119likevirus* (*Cd119virus*), *Phie125likevirus* (*E125virus*), *Phihlikevirus*
- 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.Sextaevirus [17]**
- 2015.017aB create 2 species in genus *Sextaevirus* (new) in the family *Siphoviridae*: *Staphylococcus virus Sextaev*, *Staphylococcus virus SEP9*
- 2015.017bB create a new genus in the family *Siphoviridae*, order *Caudovirales*
- 2015.017cB name the genus *Sextaevirus*
- 2015.017dB designate *Staphylococcus virus Sextaev* 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 *Tunlikevirus* (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 <i>Escherichia phage Tls</i> (proposed name <i>Escherichia virus Tls</i>) as type species of the new genus		<i>Escherichia phage Rtp</i> and <i>Escherichia phage Tls</i> from genus <i>Tunaliavirus</i>
2015.019fB	create species in genus <i>Rtpvirus</i> (new) in the family <i>Siphoviridae</i> : <i>Escherichia virus ACGM12</i>	2015.019xB	reassign species <i>Cronobacter phage Esp2949-1</i> to be unassigned in the subfamily <i>Tunavirinae</i> (new)
2015.019gB	create a new genus in the subfamily <i>Tunavirinae</i> (new), family <i>Siphoviridae</i> , order <i>Caudovirales</i>	2015.019yB	reassign species <i>Escherichia phage Tls</i> (proposed name <i>Escherichia virus Tls</i>) to the new genus <i>Tlsvirus</i> in the subfamily <i>Tunavirinae</i> (new)
2015.019hB	name the genus <i>Rtpvirus</i>	2015.019zB	reassign species <i>Enterobacter phage F20</i> (proposed name <i>Enterobacter virus F20</i>) to the new genus <i>Kp36virus</i> in the subfamily <i>Tunavirinae</i> (new)
2015.019iB	designate <i>Escherichia phage Rtp</i> (proposed name <i>Escherichia virus RTP</i>) as type species of the new genus	2015.019aaB	reassign species <i>Escherichia phage Rtp</i> (proposed name <i>Escherichia virus RTP</i>) to the new genus <i>Rtpvirus</i> in the subfamily <i>Tunavirinae</i> (new)
2015.019jB	create 2 species in genus <i>Kp36virus</i> (new) in the subfamily <i>Tunavirinae</i> (new), family <i>Siphoviridae</i> : <i>Klebsiella virus KP36</i> , <i>Klebsiella virus 1513</i>	2015.019abB	reassign species <i>Escherichia virus Jk06</i> (proposed name <i>Escherichia virus Jk06</i>) and <i>Escherichia phage Eb49</i> (proposed name <i>Escherichia phage Eb49</i>) and <i>Escherichia phage Rogue1</i> (proposed name <i>Escherichia virus Rogue1</i>) to the new genus <i>Rogue1virus</i> in the subfamily <i>Tunavirinae</i> (new)
2015.019kB	create a new genus in the subfamily <i>Tunavirinae</i> (new), family <i>Siphoviridae</i> , order <i>Caudovirales</i>		
2015.019lB	name the genus <i>Kp36virus</i>		
2015.019mB	designate <i>Klebsiella virus KP36</i> as type species of the new genus		
2015.019nB	create 5 species in genus <i>Rogue1virus</i> (new) in the subfamily <i>Tunavirinae</i> (new), family <i>Siphoviridae</i> : <i>Escherichia virus AHS24</i> , <i>Escherichia virus KP26</i> , <i>Escherichia virus AHP42</i> , <i>Escherichia virus AKS96</i> , <i>Escherichia virus E41c</i>		
2015.019oB	create a new genus in the subfamily <i>Tunavirinae</i> (new), family <i>Siphoviridae</i> , order <i>Caudovirales</i>	2015.020a-aeB.A.v4.Tevenvirinae [20]	
2015.019pB	name the genus <i>Rogue1virus</i>	2015.020aB	create 10 species in genus <i>T4likevirus</i> (proposed name <i>T4virus</i>) in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> : <i>Escherichia virus AR1</i> , <i>Escherichia virus Ime09</i> , <i>Escherichia virus E112</i> , <i>Escherichia virus ECML134</i> , <i>Escherichia virus RB3</i> , <i>Escherichia virus C40</i> , <i>Shigella virus Shf12</i> , <i>Shigella virus Pss1</i> , <i>Yersinia virus PST</i> , <i>Yersinia virus D1</i>
2015.019qB	designate <i>Escherichia phage Rogue1</i> (proposed name <i>Escherichia virus Rogue1</i>) as type species of the new genus	2015.020bB	create a new genus in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> , order <i>Caudovirales</i>
2015.019rB	create a new subfamily in the family <i>Siphoviridae</i> , order <i>Caudovirales</i>	2015.020cB	name the genus <i>Rb49virus</i>
2015.019sB	name the new subfamily <i>Tunavirinae</i>	2015.020dB	designate <i>Escherichia phage RB49</i> (existing species) to be renamed <i>Escherichia virus RB49</i> as type species of the new genus
2015.019tB	assign to the new subfamily the 5 genera, <i>Tunaliavirus</i> (existing, proposed name <i>Tl1virus</i>), <i>Rtpvirus</i> (new), <i>Tlsvirus</i> (new), <i>Kp36virus</i> (new) and <i>Rogue1virus</i> (new)	2015.020eB	create 3 species in genus <i>Rb69virus</i> (new) in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> : <i>Escherichia virus JS09</i> , <i>Escherichia virus HX01</i> , <i>Shigella virus UTAM</i>
2015.019uB	remove genus <i>Tunaliavirus</i> from its position of unassigned in the family <i>Siphoviridae</i>	2015.020fB	create a new genus in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> , order <i>Caudovirales</i>
2015.019vB	reassign the above genus to the subfamily <i>Tunavirinae</i> (new), family <i>Siphoviridae</i>		
2015.019wB	remove 7 species (<i>Cronobacter phage Esp2949-1</i> , <i>Enterobacter phage F20</i> , <i>Escherichia phage Eb49</i> , <i>Escherichia phage Jk06</i> , <i>Escherichia phage Rogue1</i> ,		

2015.020gB	name the genus <i>Rb69virus</i>	<i>Escherichia phage JS98</i> , <i>Escherichia phage phi1</i> , <i>Escherichia phage RB16</i> , <i>Escherichia phage RB32</i> , <i>Escherichia phage RB43</i> , <i>Escherichia phage RB49</i> , <i>Escherichia phage RB69</i> and <i>Pseudomonas phage 42</i>) from genus <i>T4likevirus</i>
2015.020hB	designate <i>Escherichia phage RB69</i> (existing species) to be renamed <i>Escherichia virus RB69</i> as type species of the new genus	
2015.020iB	create 4 species in genus <i>Js98virus</i> (new) in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> : <i>Escherichia virus IME08</i> , <i>Escherichia virus Bp7</i> , <i>Escherichia virus JS10</i> , <i>Escherichia virus VR5</i>	2015.020zB reassign 2 of the above species (<i>Escherichia phage RB49</i> [proposed name <i>Escherichia virus RB49</i>] and <i>Escherichia phage phi1</i> [proposed name <i>Escherichia virus phi1</i>]) to genus <i>Rb49virus</i> (new)
2015.020jB	create a new genus in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> , order <i>Caudovirales</i>	2015.020aaB reassign <i>Escherichia phage RB69</i> (proposed name <i>Escherichia virus RB69</i>) to genus <i>Rb69virus</i> (new)
2015.020kB	name the genus <i>Js98virus</i>	2015.020abB reassign <i>Escherichia phage JS98</i> (proposed name <i>Escherichia virus JS98</i>) to genus <i>JS98virus</i> (new)
2015.020lB	designate <i>Escherichia phage JS98</i> (existing species) to be renamed <i>Escherichia virus JS98</i> as type species of the new genus	2015.020acB reassign 5 of the above species (<i>Enterobacteria phage SV14</i> [proposed name <i>Enterobacteria virus SV14</i>], <i>Escherichia phage Rb16</i> [proposed name <i>Escherichia virus Rb16</i>], <i>Escherichia phage RB32</i> [proposed name <i>Escherichia virus RB32</i>], <i>Escherichia phage Rb43</i> [proposed name <i>Escherichia virus Rb43</i>] and <i>Pseudomonas phage 42</i> [proposed name <i>Pseudomonas virus 42</i>]) to be unassigned in subfamily <i>Tevenvirinae</i>
2015.020mB	create 5 species in genus <i>Sp18virus</i> (new) in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> : <i>Shigella virus SP18</i> , <i>Escherichia virus VR7</i> , <i>Escherichia virus VR20</i> , <i>Escherichia virus VR25</i> , <i>Escherichia virus VR26</i>	2015.020adB reassign <i>Aeromonas phage 25</i> (proposed name <i>Aeromonas virus 25</i>) and <i>Aeromonas phage 31</i> (proposed name <i>Aeromonas virus 31</i>) to genus <i>Secunda5virus</i> (new)
2015.020nB	create a new genus in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> , order <i>Caudovirales</i>	2015.020aeB reassign <i>Aeromonas phage 44RR2.8t</i> (proposed name <i>Aeromonas virus 44RR2</i>) to genus <i>Biquartavirus</i> (new)
2015.020oB	name the genus <i>Sp18virus</i>	
2015.020pB	designate <i>Shigella virus SP18</i> as type species of the new genus	
2015.020qB	create 2 species in genus <i>Sp16virus</i> (new) in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> : <i>Salmonella virus STML198</i> , <i>Salmonella virus S16</i>	
2015.020rB	create a new genus in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> , order <i>Caudovirales</i>	
2015.020sB	name the genus <i>Sp16virus</i>	2015.021a, bB.A.v3.T5virus_4sp1rem [21]
2015.020tB	designate <i>Salmonella virus S16</i> as type species of the new genus	2015.021aB create 4 species in genus <i>T5likevirus</i> (proposed name <i>T5virus</i>), family <i>Siphoviridae</i> : <i>Escherichia virus FFH1</i> , <i>Salmonella virus Shivani</i> , <i>Salmonella virus Stitch</i> , <i>Escherichia virus DT57C</i>
2015.020uB	create 2 species in genus <i>Cc31virus</i> (new) in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> : <i>Enterobacter virus CC31</i> , <i>Enterobacter virus PG7</i>	2015.021bB Remove (abolish) species <i>Vibrio phage 149</i> (type IV) from genus <i>T5likevirus</i>
2015.020vB	create a new genus in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> , order <i>Caudovirales</i>	
2015.020wB	name the genus <i>Cc31virus</i>	2015.022a-dB.A.v2.Tg1virus [22]
2015.020xB	designate <i>Enterobacter virus CC31</i> as type species of the new genus	2015.022aB create 2 species in genus <i>Tg1virus</i> (new) in the family <i>Myoviridae</i> : <i>Yersinia virus TGI</i> , <i>Yersinia virus RIRT</i>
2015.020yB	remove 12 species (<i>Aeromonas phage 25</i> , <i>Aeromonas phage 31</i> , <i>Aeromonas phage 44RR2.8t</i> , <i>Enterobacteria phage SV14</i> ,	2015.022bB create a new genus in the family <i>Myoviridae</i> , order <i>Caudovirales</i>

- 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.Pbunlikevirus_5sp [30]**
- 2015.030aB create 5 species in genus *Pbunlikevirus* (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*, *Haloarcula 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-dB.A.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* *ECBPI*, *Escherichia*

- virus IME11, Escherichia virus ECIUPM, 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 family *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_spre [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 *Itovirus*, 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 <i>Dyotaupapillomavirus</i> (new), family <i>Papillomaviridae</i> : <i>Dyotaupapillomavirus</i> <i>1</i>	2015.014xD	name the genus <i>Dyophipapillomavirus</i>
2015.014hD	create species in genus <i>Dyousipapillomavirus</i> (new), family <i>Papillomaviridae</i> : <i>Dyousipapillomavirus 1</i>	2015.014yD	designate <i>Dyophipapillomavirus 1</i> as the type species of the new genus
2015.014iD	create species in genus <i>Dyophipapillomavirus</i> (new), family <i>Papillomaviridae</i> : <i>Dyophipapillomavirus</i> <i>1</i>	2015.014zD	create new genus in the family <i>Papillomaviridae</i>
2015.014jD	create species in genus <i>Dyochipapillomavirus</i> (new), family <i>Papillomaviridae</i> : <i>Dyochipapillomavirus</i> <i>1</i>	2015.014aaD	name the genus <i>Dyochipapillomavirus</i>
2015.014kD	create species in genus <i>Dyopsipapillomavirus</i> (new), family <i>Papillomaviridae</i> : <i>Dyopsipapillomavirus</i> <i>1</i>	2015.014abD	designate <i>Dyochipapillomavirus 1</i> as the type species of the new genus
2015.014lD	create species in genus <i>Dyomegapapillomavirus</i> (new), family <i>Papillomaviridae</i> : <i>Dyomegapapillomavirus 1</i>	2015.014acD	create new genus in the family <i>Papillomaviridae</i>
2015.014mD	create species in genus <i>Treisidapapillomavirus</i> (new), family <i>Papillomaviridae</i> : <i>Treisidapapillomavirus 1</i>	2015.014adD	name the genus <i>Dyopsipapillomavirus</i>
2015.014nD	create species in genus <i>Treisepsilonpapillomavirus</i> (new), family <i>Papillomaviridae</i> : <i>Treisepsilonpapillomavirus 1</i>	2015.014aeD	designate <i>Dyopsipapillomavirus 1</i> as the type species of the new genus
2015.014oD	create species in genus <i>Treiszetapapillomavirus</i> (new), family <i>Papillomaviridae</i> : <i>Treiszetapapillomavirus 1</i>	2015.014afD	create new genus in the family <i>Papillomaviridae</i>
2015.014pD	create species in genus <i>Treisetapapillomavirus</i> (new), family <i>Papillomaviridae</i> : <i>Treisetapapillomavirus</i> <i>1</i>	2015.014agD	name the genus <i>Dyomegapapillomavirus</i>
2015.014qD	create new genus in the family <i>Papillomaviridae</i>	2015.014ahD	designate <i>Dyomegapapillomavirus 1</i> as the type species of the new genus
2015.014rD	name the genus <i>Taupapillomavirus</i>	2015.014aiD	create new genus in the family <i>Papillomaviridae</i>
2015.014sD	designate <i>Taupapillomavirus 1</i> as the type species of the new genus	2015.014ajD	name the genus <i>Treisidapapillomavirus</i>
2015.014tD	create new genus in the family <i>Papillomaviridae</i>	2015.014akD	designate <i>Treisidapapillomavirus 1</i> as the type species of the new genus
2015.014uD	name the genus <i>Dyousipapillomavirus</i>	2015.014alD	create new genus in the family <i>Papillomaviridae</i>
2015.014vD	designate <i>Dyousipapillomavirus 1</i> as the type species of the new genus	2015.014amD	name the genus <i>Treisepsilonpapillomavirus</i>
2015.014wD	create new genus in the family <i>Papillomaviridae</i>	2015.014anD	designate <i>Treisepsilonpapillomavirus 1</i> as the type species of the new genus
		2015.014aoD	create new genus in the family <i>Papillomaviridae</i>
		2015.014apD	name the genus <i>Treiszetapapillomavirus</i>
		2015.014aqD	designate <i>Treiszetapapillomavirus 1</i> as the type species of the new genus
		2015.014arD	create new genus in the family <i>Papillomaviridae</i>
		2015.014asD	name the genus <i>Treisetapapillomavirus</i>
		2015.014atD	designate <i>Treisetapapillomavirus 1</i> as the type species of the new genus
		2015.015a-aaD.A.v2.Polyomaviridae_rev [70]	
		2015.015aD	create 34 species in genus <i>Alphapolyomavirus</i> (new), family <i>Polyomaviridae</i> : <i>Acerodon celebensis</i> <i>polyomavirus 1</i> , <i>Artibeus planirostris</i> <i>polyomavirus 2</i> , <i>Artibeus planirostris</i> <i>polyomavirus 3</i> , <i>Ateles paniscus</i> <i>polyomavirus 1</i> , <i>Cardioderma cor</i> <i>polyomavirus 1</i> , <i>Carollia perspicillata</i> <i>polyomavirus 1</i> , <i>Chlorocebus pygerythrus</i> <i>polyomavirus 1</i> , <i>Chlorocebus pygerythrus</i> <i>polyomavirus 3</i> , <i>Dobsonia moluccensis</i> <i>polyomavirus 1</i> , <i>Eidolon helvum</i>

	<i>polyomavirus 1, Gorilla gorilla</i>		<i>polyomavirus 1, Pygoscelis adeliae</i>
	<i>polyomavirus 1, Human polyomavirus 5,</i>		<i>polyomavirus 1, Pyrrhula pyrrhula</i>
	<i>Human polyomavirus 8, Human</i>		<i>polyomavirus 1, Serinus canaria</i>
	<i>polyomavirus 9, Human polyomavirus 12,</i>		<i>polyomavirus 1</i>
	<i>Human polyomavirus 13, Macaca</i>	2015.015dD	create 4 species in genus
	<i>fascicularis polyomavirus 1, Molossus</i>		<i>Deltapolyomavirus</i> (new), family
	<i>molossus polyomavirus 1, Otomops</i>		<i>Polyomaviridae: Human polyomavirus 6,</i>
	<i>martiensseni polyomavirus 1, Otomops</i>		<i>Human polyomavirus 7, Human</i>
	<i>martiensseni polyomavirus 2, Pan</i>	2015.015eD	<i>polyomavirus 10, Human polyomavirus 11</i>
	<i>troglodytes polyomavirus 1, Pan</i>		create new genus in the family
	<i>troglodytes polyomavirus 2, Pan</i>		<i>Polyomaviridae</i>
	<i>troglodytes polyomavirus 3, Pan</i>	2015.015fD	name the genus <i>Alphapolyomavirus</i>
	<i>troglodytes polyomavirus 4, Pan</i>	2015.015gD	designate <i>Mus musculus polyomavirus 1</i> as
	<i>troglodytes polyomavirus 5, Pan</i>		the type species of the new genus
	<i>troglodytes polyomavirus 6, Pan</i>	2015.015hD	create new genus in the family
	<i>troglodytes polyomavirus 7, Papio</i>		<i>Polyomaviridae</i>
	<i>cynocephalus polyomavirus 1,</i>	2015.015iD	name the genus <i>Betapolyomavirus</i>
	<i>Piliocolobus rufomitratu polyomavirus 1,</i>	2015.015jD	designate <i>Macaca mulatta polyomavirus 1</i>
	<i>Pongo abelii polyomavirus 1, Pongo</i>		as the type species of the new genus
	<i>pygmaeus polyomavirus 1, Procyon lotor</i>	2015.015kD	create new genus in the family
	<i>polyomavirus 1, Pteropus vampyrus</i>		<i>Polyomaviridae</i>
	<i>polyomavirus 1, Sturnira lilium</i>	2015.015lD	name the genus <i>Gammapolyomavirus</i>
	<i>polyomavirus 1</i>	2015.015mD	designate <i>Aves polyomavirus 1</i> as the type
2015.015bD	create 22 species in genus		species of the new genus
	<i>Betapolyomavirus</i> (new), family	2015.015nD	create new genus in the family
	<i>Polyomaviridae: Acerodon celebensis</i>		<i>Polyomaviridae</i>
	<i>polyomavirus 2, Artibeus planirostris</i>	2015.015oD	name the genus <i>Deltapolyomavirus</i>
	<i>polyomavirus 1, Cebus albifrons</i>	2015.015pD	designate <i>Human polyomavirus 6</i> as the
	<i>polyomavirus 1, Cercopithecus erythrotis</i>		type species of the new genus
	<i>polyomavirus 1, Chlorocebus pygerythrus</i>	2015.015qD	remove the genus <i>Polyomavirus</i>
	<i>polyomavirus 2, Desmodus rotundus</i>	2015.015rD	remove 4 species (<i>African green monkey</i>
	<i>polyomavirus 1, Dobsonia moluccensis</i>		<i>polyomavirus, Simian virus 12, Baboon</i>
	<i>polyomavirus 2, Dobsonia moluccensis</i>		<i>polyomavirus 2, Rabbit kidney vacuolating</i>
	<i>polyomavirus 3, Equus caballus</i>		<i>virus</i>) from their unassigned position in the
	<i>polyomavirus 1, Human polyomavirus 3,</i>	2015.015sD	family <i>Polyomaviridae</i>
	<i>Human polyomavirus 4, Loxodonta</i>		remove species <i>Murine polyomavirus</i> and
	<i>africana polyomavirus 1, Mastomys</i>		<i>Hamster polyomavirus</i> from genus
	<i>natalensis polyomavirus 1, Meles meles</i>		<i>Polyomavirus</i> (to be removed) in the
	<i>polyomavirus 1, Miniopterus africanus</i>	2015.015tD	family <i>Polyomaviridae</i>
	<i>polyomavirus 1, Myotis lucifugus</i>		reassign the above species to genus
	<i>polyomavirus 1, Papio cynocephalus</i>	2015.015uD	<i>Alphapolyomavirus</i> (new), family
	<i>polyomavirus 2, Pteronotus davyi</i>		<i>Polyomaviridae</i>
	<i>polyomavirus 1, Pteronotus parnellii</i>		remove 4 species (<i>BK polyomavirus, JC</i>
	<i>polyomavirus 1, Saimiri boliviensis</i>		<i>polyomavirus, Murine pneumotropic virus,</i>
	<i>polyomavirus 1, Saimiri sciureus</i>		<i>Simian virus 40</i>) from genus <i>Polyomavirus</i>
	<i>polyomavirus 1, Zalophus californianus</i>		(to be removed) in the family
	<i>polyomavirus 1</i>	2015.015vD	<i>Polyomaviridae</i>
2015.015cD	create 6 species in genus		reassign the above species to genus
	<i>Gammapolyomavirus</i> (new), family		<i>Betapolyomavirus</i> (new), family
	<i>Polyomaviridae: Anser anser</i>		<i>Polyomaviridae</i>
	<i>polyomavirus 1, Corvus monedula</i>	2015.015wD	remove species <i>Budgerigar fledgling</i>
	<i>polyomavirus 1, Cracticus torquatus</i>		<i>disease polyomavirus</i> from genus

- 2015.015xD *Polyomavirus* (to be removed) in the family *Polyomaviridae*
reassign the above species to genus *Gammampolyomavirus* (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*: *Suca 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
- 2015.001kF create a new genus in the family *Lavidaviridae* (new)
- 2015.001kF name the new genus *Mavirus*
- 2015.001kF designate *Cafeteriavirus-dependent mavirus* as type species in the new genus
- 2015.001kF create a new unassigned family
- 2015.001kF name the new family *Lavidaviridae*
- 2015.001kF 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*

- 2015.007mP *virus, Japanese iris necrotic ring virus and Turnip crinkle virus*) from the genus *Carmovirus*, family *Tombusviridae*
- 2015.007nP reassign the above 4 species to the genus *Betacarmovirus* (new), family *Tombusviridae*
- 2015.007oP 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*
- 2015.007pP reassign the above species to the genus *Gammacarmovirus* (new), family *Tombusviridae*
- 2015.007qP remove 4 species (*Ahlum waterborne virus, Bean mild mosaic virus, Cucumber soil-borne virus and Weddel waterborne virus* from the genus *Carmovirus*, family *Tombusviridae*
- 2015.007rP 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.Sarothroviridae [123]

- 2015.002aS create species in genus *Macronovirus* (new), family *Sarothroviridae* (new):
Macrobrachium satellite virus 1
- 2015.002bS create new genus in the family *Sarothroviridae* (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 *Sarothroviridae*
- 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 *Nidovirales*: *Coronavirus HKU15, White-eye coronavirus HKU16, Night heron coronavirus HKU19, Wigeon coronavirus HKU20, Common moorhen coronavirus HKU21*
- 2015.003dS create species in genus *Bafnavirus*, 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 *Sinaivirus*
- 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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 26. Krupovic M, Kropinski AM, Adriaenssens EM (2015) ICTV taxonomic proposal 2015.026a-rB.A.v3.Bullavirinae. In family *Microviridae*, create 3 genera (*Alpha3microvirus*, *Phix174microvirus* and *G4microvirus*), including 9 new species and 5 existing species reassigned from genus *Microvirus*. Assign the new genera to a new subfamily *Bullavirinae*. <http://www.ictvonline.org/proposals-15/2015.026a-rB.A.v3.Bullavirinae.pdf>
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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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