

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 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*
- 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 *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. *Andromedallikevirus* to *Andromedavirus*). Also make the following individual name changes (proposed name in brackets):
3alikevirus (*Triavirus*), *77likevirus* (*Biseptimavirus*), *Ahjdlikevirus* (*P68virus*), *Bppunallikevirus* (*Bpp1virus*), *Cjwunallikevirus* (*Cjw1virus*), *Cp8unallikevirus* (*Cp8virus*), *Felixounallikevirus* (*Felixo1virus*), *Halolikevirus* (*Liefievirus*), *Hapunallikevirus* (*Hapunavirus*), *Hpunallikevirus* (*Hp1virus*), *I3likevirus* (*Bxz1virus*), *Iebhlikevirus* (*Cecivirus*), *Pbiunallikevirus* (*Pbi1virus*), *Pbunallikevirus* (*Pbunavirus*), *Pgonellikevirus* (*Pg1virus*), *Phic3unallikevirus* (*Phic31virus*), *Phicd119likevirus* (*Cd119virus*), *Phie125likevirus* (*E125virus*), *Phihlikevirus* (*Myohalovirus*), *Phijlunallikevirus* (*Phijl1virus*), *Psimunallikevirus* (*Psimunavirus*), *Punallikevirus* (*P1virus*), *Sfi1unallikevirus* (*Sfi11virus*), *Sfi21dtunallikevirus* (*Sfi21dt1virus*), *Skunallikevirus* (*Sk1virus*), *Spounallikevirus* (*Spo1virus*), *Tp2unallikevirus* (*Tp21virus*), *Tunallikevirus* (*T1virus*), *Viunallikevirus* (*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*

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| 2015.011bB | create a new genus in the family <i>Siphoviridae</i> , order <i>Caudovirales</i> |
| 2015.011cB | name the genus <i>P70virus</i> |
| 2015.011dB | designate <i>Listeria virus P70</i> as type species of the new genus |
| 2015.012a-eB.A.v3.P100virus [12] | |
| 2015.012aB | create a new genus in the subfamily <i>Spounavirinae</i> , family <i>Myoviridae</i> , order <i>Caudovirales</i> |
| 2015.012bB | name the genus <i>P100virus</i> |
| 2015.012cB | designate <i>Listeria virus P100</i> as type species of the new genus |
| 2015.012dB | remove species <i>Listera phage P100</i> and <i>Listera phage A511</i> from genus <i>Twortlikevirus</i> |
| 2015.012eB | reassign the above two species to genus <i>P100virus</i> |
| 2015.013a-cB.A.v3.Biquartavirus [13] | |
| 2015.013aB | create a new genus in the family <i>Myoviridae</i> , order <i>Caudovirales</i> |
| 2015.013bB | name the genus <i>Biquartavirus</i> |
| 2015.013cB | designate <i>Aeromonas virus 44RR2</i> as type species of the new genus |
| 2015.014aB.A.v3.Phieco32virus_4sp [14] | |
| 2015.014aB | create 4 species within the genus <i>Phieco32virus</i> , family <i>Podoviridae</i> : <i>Escherichia virus SU10</i> , <i>Escherichia virus NJ01</i> , <i>Escherichia virus ECB2</i> , <i>Escherichia virus Septima11</i> |
| 2015.015a-dB.A.v2.Psavirus [15] | |
| 2015.015aB | create 2 species in genus <i>Psavirus</i> (new) in the family <i>Siphoviridae</i> : <i>Listeria virus PSA</i> , <i>Listeria virus LP302</i> |
| 2015.015bB | create a new genus in the family <i>Siphoviridae</i> , order <i>Caudovirales</i> |
| 2015.015cB | name the genus <i>Psavirus</i> |
| 2015.015dB | designate <i>Listeria virus PSA</i> as type species of the new genus |
| 2015.016a-dB.A.v3.Secunda5virus [16] | |
| 2015.016aB | create 4 species in genus <i>Secunda5virus</i> (new), family <i>Myoviridae</i> : <i>Aeromonas virus AS4</i> , <i>Stenotrophomonas virus IME13</i> , <i>Aeromonas virus Aes12</i> , <i>Aeromonas virus Aes508</i> |
| 2015.016bB | create a new genus in the family <i>Myoviridae</i> , order <i>Caudovirales</i> |
| 2015.016cB | name the genus <i>Secunda5virus</i> |
| 2015.016dB | designate <i>Aeromonas virus 25</i> (see 2015.020adB) as type species of the new genus |
| 2015.017a-dB.A.v3.Sextaecvirus [17] | |
| 2015.017aB | create 2 species in genus <i>Sextaecvirus</i> (new) in the family <i>Siphoviridae</i> : <i>Staphylococcus virus Sextaec</i> , <i>Staphylococcus virus SEP9</i> |
| 2015.017bB | create a new genus in the family <i>Siphoviridae</i> , order <i>Caudovirales</i> |
| 2015.017cB | name the genus <i>Sextaecvirus</i> |
| 2015.017dB | designate <i>Staphylococcus virus Sextaec</i> as type species of the new genus |
| 2015.018a-dB.A.v2.Silviavirus [18] | |
| 2015.018aB | create 2 species in genus <i>Silviavirus</i> (new), family <i>Myoviridae</i> : <i>Staphylococcus virus Remus</i> , <i>Staphylococcus virus SA11</i> |
| 2015.018bB | create a new genus in the family <i>Myoviridae</i> , order <i>Caudovirales</i> |
| 2015.018cB | name the genus <i>Silviavirus</i> |
| 2015.018dB | designate <i>Staphylococcus virus Remus</i> as type species of the new genus |
| 2015.019a-abB.A.v3.Tunavirinae [19] | |
| 2015.019aB | create 2 species in genus <i>Tunlikevirus</i> (proposed name <i>Tlivirus</i>) in the family <i>Siphoviridae</i> |
| 2015.019bB | create 2 species in genus <i>Tlsvirus</i> (new) in the subfamily <i>Tunavirinae</i> (new), family <i>Siphoviridae</i> : <i>Escherichia virus ADB2</i> , <i>Shigella virus PSf2</i> , <i>Salmonella virus SP126</i> , <i>Citrobacter virus Stevie</i> |
| 2015.019cB | create a new genus in the subfamily <i>Tunavirinae</i> (new), family <i>Siphoviridae</i> , order <i>Caudovirales</i> |
| 2015.019dB | name the genus <i>Tlsvirus</i> |
| 2015.019eB | designate <i>Escherichia phage Tls</i> (proposed name <i>Escherichia virus Tls</i>) as type species of the new genus |

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| 2015.019fB | create species in genus <i>Rtpvirus</i> (new) in the family <i>Siphoviridae</i> : <i>Escherichia virus ACGM12</i> |
| 2015.019gB | create a new genus in the subfamily <i>Tunavirinae</i> (new), family <i>Siphoviridae</i> , order <i>Caudovirales</i> |
| 2015.019hB | name the genus <i>Rtpvirus</i> |
| 2015.019iB | designate <i>Escherichia phage Rtp</i> (proposed name <i>Escherichia virus RTP</i>) as type species of the new genus |
| 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> |
| 2015.019kB | 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> (proposed name <i>Escherichia virus RTP</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> |
| 2015.019pB | name the genus <i>Rogue1virus</i> |
| 2015.019qB | designate <i>Escherichia phage Rogue1</i> (proposed name <i>Escherichia virus Rogue1</i>) as type species of the new genus |
| 2015.019rB | create a new subfamily in the family <i>Siphoviridae</i> , order <i>Caudovirales</i> |
| 2015.019sB | name the new subfamily <i>Tunavirinae</i> |
| 2015.019tB | assign to the new subfamily the 5 genera, <i>Tunalikevirus</i> (existing, proposed name <i>Tl1virus</i>), <i>Rtpvirus</i> (new), <i>Tlsvirus</i> (new), <i>Kp36virus</i> (new) and <i>Rogue1virus</i> (new) |
| 2015.019uB | remove genus <i>Tunalikevirus</i> from its position of unassigned in the family <i>Siphoviridae</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> , <i>Escherichia phage Rtp</i> and <i>Escherichia phage Tls</i>) from genus <i>Tunalikevirus</i> |
| 2015.019xB | reassign species <i>Cronobacter phage Esp2949-1</i> to be unassigned in the subfamily <i>Tunavirinae</i> (new) |
| 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) |
| 2015.019zB | 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) |
| 2015.019aaB | 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) |
| 2015.019abB | 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) |
| 2015.020a-aeB.A.v4.Tevenvirinae [20] | |
| 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> |
| 2015.020bB | create a new genus in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> , order <i>Caudovirales</i> |
| 2015.020cB | name the genus <i>Rb49virus</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 |

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| 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> |
| 2015.020fB | create a new genus in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> , order <i>Caudovirales</i> |
| 2015.020gB | name the genus <i>Rb69virus</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> |
| 2015.020jB | create a new genus in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> , order <i>Caudovirales</i> |
| 2015.020kB | name the genus <i>Js98virus</i> |
| 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 |
| 2015.020mB | create 5 species in genus <i>Sp18virus</i> (new) in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> : <i>Shigella virus SPI8</i> , <i>Escherichia virus VR7</i> , <i>Escherichia virus VR20</i> , <i>Escherichia virus VR25</i> , <i>Escherichia virus VR26</i> |
| 2015.020nB | create a new genus in the subfamily <i>Tevenvirinae</i> (new), family <i>Myoviridae</i> , order <i>Caudovirales</i> |
| 2015.020oB | name the genus <i>Sp18virus</i> |
| 2015.020pB | designate <i>Shigella virus SPI8</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> |
| 2015.020tB | designate <i>Salmonella virus S16</i> as type species of the new genus |
| 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> |
| 2015.020vB | 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> |
| 2015.020xB | designate <i>Enterobacter virus CC31</i> as type species of the new genus |
| 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> , <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> |
| 2015.020zB | 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) |
| 2015.020aaB | reassign <i>Escherichia phage RB69</i> (proposed name <i>Escherichia virus RB69</i>) to genus <i>Rb69virus</i> (new) |
| 2015.020abB | reassign <i>Escherichia phage JS98</i> (proposed name <i>Escherichia virus JS98</i>) to genus <i>JS98virus</i> (new) |
| 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> |

- 2015.020adB 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 TGI*, *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 TGI* 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]**
- 2015.025dB 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 *Sp3Ivirus* (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 *Sp3Ivirus*
- 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*

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| 2015.044iB | designate <i>Escherichia virus K1G</i> as type species of the new genus |
| 2015.044jB | create a new subfamily in the family <i>Siphoviridae</i> , order <i>Caudovirales</i> |
| 2015.044kB | name the subfamily <i>Guernseyvirinae</i> |
| 2015.044lB | assign 3 genera [<i>Jerseylikevirus</i> (proposed name <i>Jerseyvirus</i>), <i>Sp31virus</i> (new) and <i>K1gvirus</i> (new)] to the new subfamily |
| 2015.044mB | remove genus <i>Jerseylikevirus</i> from its position of unassigned in the family <i>Siphoviridae</i> |
| 2015.044nB | reassign the above genus to subfamily <i>Guernseyvirinae</i> (new), family <i>Siphoviridae</i> |
| 2015.045a-dB.A.v2.Rheph4virus [45] | |
| 2015.045aB | create species <i>Rhizobium virus RHEph4</i> in genus <i>Rheph4virus</i> (new) in the family <i>Myoviridae</i> : <i>Rhizobium virus RHEph4</i> |
| 2015.045bB | create a new genus in the family <i>Myoviridae</i> , order <i>Caudovirales</i> |
| 2015.045cB | name the genus <i>Rheph4virus</i> |
| 2015.045dB | designate <i>Rhizobium virus RHEph4</i> as type species of the new genus |
| 2015.047aB.A.v2.Felixo1virus_7sp [46] | |
| 2015.047aB | create 7 species in genus <i>Felixounalikevirus</i> (proposed name <i>Felixo1virus</i>), family <i>Myoviridae</i> , order <i>Caudovirales</i> : <i>Salmonella virus UAB87</i> , <i>Escherichia virus VpaE1</i> , <i>Salmonella virus Mushroom</i> , <i>Escherichia virus JH2</i> , <i>Escherichia virus AYO145A</i> , <i>Salmonella virus HB2014</i> , <i>Escherichia virus EC6</i> |
| 2015.048a-dB.A.v2.Ssp2virus [47] | |
| 2015.048aB | create 2 species in the genus <i>Ssp2virus</i> (new), family <i>Siphoviridae</i> : <i>Vibrio virus SSP002</i> , <i>Vibrio virus MAR10</i> |
| 2015.048bB | create a new genus in the family <i>Siphoviridae</i> , order <i>Caudovirales</i> |
| 2015.048cB | name the genus <i>Ssp2virus</i> |
| 2015.048dB | designate <i>Vibrio virus SSP002</i> as type species of the new genus |
| 2015.049a-dB.A.v2.Cvm10virus [48] | |
| 2015.049aB | create 2 species in genus <i>Cvm10virus</i> (new) in the family <i>Myoviridae</i> : <i>Escherichia virus CVM10</i> , <i>Escherichia virus ep3</i> |
| 2015.049bB | create a new genus in the family <i>Myoviridae</i> , order <i>Caudovirales</i> |
| 2015.049cB | name the genus <i>Cvm10virus</i> |
| 2015.049dB | designate <i>Escherichia virus CVM10</i> as type species of the new genus |
| 2015.050a-dB.A.v2.G7cvirus [49] | |
| 2015.050aB | create 8 species in genus <i>G7cvirus</i> (new) in the family <i>Podoviridae</i> : <i>Escherichia virus G7C</i> , <i>Escherichia virus APEC7</i> , <i>Escherichia virus APEC5</i> , <i>Escherichia virus Bp4</i> , <i>Escherichia virus ECBP1</i> , <i>Escherichia virus IME11</i> , <i>Escherichia virus ECIUPM</i> , <i>Shigella virus Sb1</i> |
| 2015.050bB | create a new genus in the family <i>Podoviridae</i> , order <i>Caudovirales</i> |
| 2015.050cB | name the genus <i>G7cvirus</i> |
| 2015.050dB | designate <i>Escherichia virus G7C</i> as type species of the new genus |
| 2015.051a-dB.A.v2.Nonagvirus [50] | |
| 2015.051aB | create 4 species in genus <i>Nonagvirus</i> (new) in the family <i>Siphoviridae</i> : <i>Escherichia virus 9g</i> , <i>Escherichia virus JenK1</i> , <i>Escherichia virus JenP2</i> , <i>Escherichia virus JenP1</i> |
| 2015.051bB | create a new genus in the family <i>Siphoviridae</i> , order <i>Caudovirales</i> |
| 2015.051cB | name the genus <i>Nonagvirus</i> |
| 2015.051dB | designate <i>Escherichia virus 9g</i> as type species of the new genus |
| 2015.052aB.A.v3.Schizot4virus_sp [51] | |
| 2015.052aB | create species in genus <i>Schizot4likevirus</i> (proposed name <i>Schizot4virus</i>), subfamily <i>Tevenvirinae</i> , family <i>Myoviridae</i> , order <i>Caudovirales</i> : <i>Vibrio virus ValKK3</i> |
| 2015.053a-dB.A.v2.Seuratvirus [52] | |
| 2015.053aB | create 2 species in genus <i>Seuratvirus</i> (new) in the family <i>Siphoviridae</i> : <i>Escherichia virus Seurat</i> , <i>Escherichia virus CAjan</i> |
| 2015.053bB | create a new genus in the family <i>Siphoviridae</i> , order <i>Caudovirales</i> |
| 2015.053cB | name the genus <i>Seuratvirus</i> |
| 2015.053dB | designate <i>Escherichia virus Seurat</i> as type species of the new genus |
| 2015.054a-dB.A.v3.Septima3virus [53] | |

- 2015.054aB create 5 species in genus *Septima3virus* (new) in the family *Siphoviridae*:
Pseudomonas virus 73, *Pseudomonas virus Kakheta25*, *Pseudomonas virus Ab26*,
Burkholderia virus KLI
- 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

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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 unassigned genus
- 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_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 *Itoivirus*, 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 *Dyousilonpapillomavirus* (new), family *Papillomaviridae*: *Dyousilonpapillomavirus 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 *Dyomegapapillomavirus* (new), family *Papillomaviridae*: *Dyomegapapillomavirus 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 *Dyousipapillomavirus*

2015.014vD designate *Dyousipapillomavirus 1* as the type species of the new genus

2015.014wD create new genus in the family *Papillomaviridae*

2015.014xD name the genus *Dyophilapapillomavirus*

2015.014yD designate *Dyophilapapillomavirus 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 *Dyomegapapillomavirus*

2015.014ahD designate *Dyomegapapillomavirus 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 <i>Betapolyomavirus</i> (new), family <i>Polyomaviridae</i> : <i>Acerodon celebensis polyomavirus</i> 2, <i>Artibeus planirostris polyomavirus</i> 1, <i>Cebus albifrons polyomavirus</i> 1, <i>Cercopithecus erythrotis polyomavirus</i> 1, <i>Chlorocebus pygerythrus polyomavirus</i> 2, <i>Desmodus rotundus polyomavirus</i> 1, <i>Dobsonia moluccensis polyomavirus</i> 2, <i>Dobsonia moluccensis polyomavirus</i> 3, <i>Equus caballus polyomavirus</i> 1, <i>Human polyomavirus</i> 3, <i>Human polyomavirus</i> 4, <i>Loxodonta africana polyomavirus</i> 1, <i>Mastomys natalensis polyomavirus</i> 1, <i>Meles meles polyomavirus</i> 1, <i>Miniopterus africanus polyomavirus</i> 1, <i>Myotis lucifugus polyomavirus</i> 1, <i>Papio cynocephalus polyomavirus</i> 2, <i>Pteronotus davyi polyomavirus</i> 1, <i>Pteronotus parnellii polyomavirus</i> 1, <i>Saimiri boliviensis polyomavirus</i> 1, <i>Saimiri sciureus polyomavirus</i> 1, <i>Zalophus californianus polyomavirus</i> 1 |
| 2015.015cD | create 6 species in genus <i>Gammampolyomavirus</i> (new), family <i>Polyomaviridae</i> : <i>Anser anser polyomavirus</i> 1, <i>Corvus monedula polyomavirus</i> 1, <i>Cracticus torquatus polyomavirus</i> 1, <i>Pygoscelis adeliae polyomavirus</i> 1, <i>Pyrrhula pyrrhula polyomavirus</i> 1, <i>Serinus canaria polyomavirus</i> 1 |
| 2015.015dD | create 4 species in genus <i>Deltapolyomavirus</i> (new), family <i>Polyomaviridae</i> : <i>Human polyomavirus</i> 6, <i>Human polyomavirus</i> 7, <i>Human polyomavirus</i> 10, <i>Human polyomavirus</i> 11 |
| 2015.015eD | create new genus in the family <i>Polyomaviridae</i> |
| 2015.015fD | name the genus <i>Alphapolyomavirus</i> |
| 2015.015gD | designate <i>Mus musculus polyomavirus</i> 1 as the type species of the new genus |
| 2015.015hD | create new genus in the family <i>Polyomaviridae</i> |
| 2015.015iD | name the genus <i>Betapolyomavirus</i> |
| 2015.015jD | designate <i>Macaca mulatta polyomavirus</i> 1 as the type species of the new genus |
| 2015.015kD | create new genus in the family <i>Polyomaviridae</i> |
| 2015.015lD | name the genus <i>Gammampolyomavirus</i> |
| 2015.015mD | designate <i>Aves polyomavirus</i> 1 as the type species of the new genus |
| 2015.015nD | create new genus in the family <i>Polyomaviridae</i> |
| 2015.015oD | name the genus <i>Deltapolyomavirus</i> |
| 2015.015pD | designate <i>Human polyomavirus</i> 6 as the type species of the new genus |
| 2015.015qD | remove the genus <i>Polyomavirus</i> |
| 2015.015rD | remove 4 species (<i>African green monkey polyomavirus</i> , <i>Simian virus</i> 12, <i>Baboon polyomavirus</i> 2, <i>Rabbit kidney vacuolating virus</i>) from their unassigned position in the family <i>Polyomaviridae</i> |
| 2015.015sD | remove species <i>Murine polyomavirus</i> and <i>Hamster polyomavirus</i> from genus <i>Polyomavirus</i> (to be removed) in the family <i>Polyomaviridae</i> |
| 2015.015tD | reassign the above species to genus <i>Alphapolyomavirus</i> (new), family <i>Polyomaviridae</i> |
| 2015.015uD | remove 4 species (<i>BK polyomavirus</i> , <i>JC polyomavirus</i> , <i>Murine pneumotropic virus</i> , <i>Simian virus</i> 40) from genus <i>Polyomavirus</i> (to be removed) in the family <i>Polyomaviridae</i> |
| 2015.015vD | reassign the above species to genus <i>Betapolyomavirus</i> (new), family <i>Polyomaviridae</i> |
| 2015.015wD | remove species <i>Budgerigar fledgling disease polyomavirus</i> from genus <i>Polyomavirus</i> (to be removed) in the family <i>Polyomaviridae</i> |
| 2015.015xD | reassign the above species to genus <i>Gammampolyomavirus</i> (new), family <i>Polyomaviridae</i> |
| 2015.015yD | remove species <i>Bovine polyomavirus</i> from genus <i>Polyomavirus</i> (to be removed) in the family <i>Polyomaviridae</i> |
| 2015.015zD | reassign the above species to genus <i>Deltapolyomavirus</i> (new), family <i>Polyomaviridae</i> |
| 2015.015aaD | change the names of the following 8 species: <i>BK polyomavirus</i> to <i>Human polyomavirus</i> 1; <i>Bovine polyomavirus</i> to <i>Bos taurus polyomavirus</i> 1; <i>Budgerigar fledgling disease polyomavirus</i> to <i>Aves polyomavirus</i> 1; <i>Hamster polyomavirus</i> to <i>Mesocricetus auratus polyomavirus</i> 1; <i>JC polyomavirus</i> to <i>Human polyomavirus</i> 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

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 an unassigned genus *Dichorhavirus* (new): *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_sprent [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_sprent [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 genus 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*,
Paramyxovirinae, *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

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| 2015.007jP | remove 7 species (<i>Angelonia flower break virus</i> , <i>Calibrachoa mottle virus</i> , <i>Carnation mottle virus</i> , <i>Honeysuckle ringspot virus</i> , <i>Nootka lupine vein clearing virus</i> , <i>Pelargonium flower break virus</i> and <i>Saguaro cactus virus</i>) from the genus <i>Carmovirus</i> , family <i>Tombusviridae</i> |
| 2015.007kP | reassign the above 7 species to the genus <i>Alphacarmovirus</i> (new), family <i>Tombusviridae</i> |
| 2015.007lP | remove 4 species (<i>Cardamine chlorotic fleck virus</i> , <i>Hibiscus chlorotic ringspot virus</i> , <i>Japanese iris necrotic ring virus</i> and <i>Turnip crinkle virus</i>) from the genus <i>Carmovirus</i> , family <i>Tombusviridae</i> |
| 2015.007mP | reassign the above 4 species to the genus <i>Betacarmovirus</i> (new), family <i>Tombusviridae</i> |
| 2015.007nP | remove 4 species (<i>Cowpea mottle virus</i> , <i>Melon necrotic spot virus</i> , <i>Soybean yellow mottle mosaic virus</i> and <i>Pea stem necrosis virus</i> from the genus <i>Carmovirus</i> , family <i>Tombusviridae</i> |
| 2015.007oP | reassign the above species to the genus <i>Gammacarmovirus</i> (new), family <i>Tombusviridae</i> |
| 2015.007pP | remove 4 species (<i>Ahlum waterborne virus</i> , <i>Bean mild mosaic virus</i> , <i>Cucumber soil-borne virus</i> and <i>Weddel waterborne virus</i> from the genus <i>Carmovirus</i> , family <i>Tombusviridae</i> |
| 2015.007qP | reassign the above species as unassigned in the family <i>Tombusviridae</i> |
| 2015.007rP | remove genus <i>Carmovirus</i> from the family <i>Tombusviridae</i> |
| 2015.008aP.A.v1.Aureusvirus_sp [104] | |
| 2015.008aP | create species in genus <i>Aureusvirus</i> , family <i>Tombusviridae</i> : <i>Yam spherical virus</i> |
| 2015.009a-pP.A.v3.Plant_sat_4unassgen [105] | |
| 2015.009aP | create 3 species in the unassigned genus <i>Albetovirus</i> (new): <i>Tobacco albetovirus 1</i> , <i>Tobacco albetovirus 2</i> , <i>Tobacco albetovirus 3</i> |
| 2015.009bP | create species in the unassigned genus <i>Aumaivirus</i> (new): <i>Maize aumaivirus 1</i> |
| 2015.009cP | create species in the unassigned genus <i>Papanivirus</i> (new): <i>Panicum papanivirus 1</i> |
| 2015.009dP | create species in the unassigned genus <i>Virtovirus</i> (new): <i>Tobacco virtovirus 1</i> |
| 2015.009eP | create a new unassigned genus |
| 2015.009fP | name the genus <i>Albetovirus</i> |
| 2015.009gP | designate <i>Tobacco albetovirus 1</i> as type species of the new genus |
| 2015.009hP | create a new unassigned genus |
| 2015.009iP | name the genus <i>Aumaivirus</i> |
| 2015.009jP | designate <i>Maize aumaivirus 1</i> as type species of the new genus |
| 2015.009kP | create a new unassigned genus |
| 2015.009lP | name the genus <i>Papanivirus</i> |
| 2015.009mP | designate <i>Panicum papanivirus 1</i> as type species of the new genus |
| 2015.009nP | create a new unassigned genus |
| 2015.009oP | name the genus <i>Virtovirus</i> |
| 2015.009pP | designate <i>Tobacco virtovirus 1</i> as type species of the new genus |
| 2015.010aP.A.v1.Nanovirus_2sp [106] | |
| 2015.010aP | create two species in the genus <i>Nanovirus</i> , family <i>Nanoviridae</i> : <i>Black medic leaf roll virus</i> , <i>Pea yellow stunt virus</i> |
| 2015.011a-adP.A.v2.Betaflexiviridae_rev [107] | |
| 2015.011aP | create two species in the genus <i>Chordovirus</i> (new), subfamily <i>Trivirinae</i> (new), family <i>Betaflexiviridae</i> , order <i>Tymovirales</i> : <i>Carrot Ch virus 1</i> , <i>Carrot Ch virus 2</i> |
| 2015.011bP | create two species in the genus <i>Prunevirus</i> (new), subfamily <i>Trivirinae</i> (new), family <i>Betaflexiviridae</i> : <i>Apricot vein clearing associated virus</i> , <i>Caucasus prunus virus</i> |
| 2015.011cP | create species in the genus <i>Tepovirus</i> , subfamily <i>Trivirinae</i> (new), family <i>Betaflexiviridae</i> : <i>Prunus virus T</i> |
| 2015.011dP | create two species in the genus <i>Carlavirus</i> , subfamily <i>Quinvirinae</i> (new), family <i>Betaflexiviridae</i> : <i>Gaillardia latent virus</i> , <i>Potato virus H</i> |

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| 2015.011eP | create two species in the genus <i>Robigovirus</i> (new), subfamily <i>Quinvirinae</i> (new), family <i>Betaflexiviridae</i> : <i>Cherry rusty mottle associated virus</i> , <i>Cherry twisted leaf associated virus</i> |
| 2015.011fP | create a new genus in the subfamily <i>Trivirinae</i> (new), family <i>Betaflexiviridae</i> , order <i>Tymovirales</i> |
| 2015.011gP | name the genus <i>Chordovirus</i> |
| 2015.011hP | designate <i>Carrot Ch virus 1</i> as type species of the new genus |
| 2015.011iP | create a new genus in the subfamily <i>Trivirinae</i> (new), family <i>Betaflexiviridae</i> , order <i>Tymovirales</i> |
| 2015.011jP | name the genus <i>Divavirus</i> |
| 2015.011kP | designate <i>Diuris virus A</i> as type species of the new genus |
| 2015.011lP | create a new genus in the subfamily <i>Trivirinae</i> (new), family <i>Betaflexiviridae</i> , order <i>Tymovirales</i> |
| 2015.011mP | name the genus <i>Prunevirus</i> |
| 2015.011nP | designate <i>Apricot vein clearing associated virus</i> as type species of the new genus |
| 2015.011oP | create a new genus in the subfamily <i>Quinvirinae</i> (new), family <i>Betaflexiviridae</i> , order <i>Tymovirales</i> |
| 2015.011pP | name the genus <i>Robigovirus</i> |
| 2015.011qP | designate <i>Cherry necrotic rusty mottle virus</i> as type species of the new genus |
| 2015.011rP | create a new subfamily within the family <i>Betaflexiviridae</i> , order <i>Tymovirales</i> |
| 2015.011sP | name the subfamily <i>Trivirinae</i> |
| 2015.011tP | assign the following 8 genera to the new subfamily: <i>Capillovirus</i> , <i>Citrivirus</i> , <i>Tepovirus</i> , <i>Trichovirus</i> , <i>Vitivirus</i> , <i>Chordovirus</i> (new), <i>Divavirus</i> (new) and <i>Prunevirus</i> (new) |
| 2015.011uP | create a new subfamily within the family <i>Betaflexiviridae</i> , order <i>Tymovirales</i> |
| 2015.011vP | name the subfamily <i>Quinvirinae</i> |
| 2015.011wP | assign the following 3 genera to the new subfamily: <i>Carlavirus</i> , <i>Foveavirus</i> and <i>Robigovirus</i> (new) |
| 2015.011xP | remove 3 species (<i>Diuris virus A</i> , <i>Diuris virus B</i> and <i>Hardenbergia virus A</i>) from their present, unassigned, position in the family <i>Betaflexiviridae</i> |
| 2015.011yP | reassign the above 3 species to genus <i>Divavirus</i> (new), subfamily <i>Trivirinae</i> (new), family <i>Betaflexiviridae</i> , order <i>Tymovirales</i> |
| 2015.011zP | remove 3 species (<i>African oil palm ringspot virus</i> , <i>Cherry green ring mottle virus</i> and <i>Cherry necrotic rusty mottle virus</i>) from their present, unassigned, position in the family <i>Betaflexiviridae</i> |
| 2015.011aaP | reassign the above 3 species to genus <i>Robigovirus</i> (new), subfamily <i>Quinvirinae</i> (new), family <i>Betaflexiviridae</i> , order <i>Tymovirales</i> |
| 2015.011abP | remove 3 species (<i>Banana mild mosaic virus</i> , <i>Banana virus X</i> and <i>Sugarcane striate mosaic-associated virus</i>) from their present, unassigned, position in the family <i>Betaflexiviridae</i> |
| 2015.011acP | reassign the above 3 species to be unassigned in the subfamily <i>Quinvirinae</i> (new), family <i>Betaflexiviridae</i> , order <i>Tymovirales</i> |
| 2015.011adP | remove (abolish) 7 species (<i>Dandelion latent virus</i> , <i>Elderberry symptomless virus</i> , <i>Honeysuckle latent virus</i> , <i>Hydrangea latent virus</i> , <i>Lilac mottle virus</i> , <i>Mulberry latent virus</i> and <i>Muskmelon vein necrosis virus</i>) from the genus <i>Carlavirus</i> , family <i>Betaflexiviridae</i> |
| 2015.012a-dP.A.v5.Platypuvirus [108] | |
| 2015.012aP | create species in the genus <i>Platypuvirus</i> (new), family <i>Alphaflexiviridae</i> , order <i>Tymovirales</i> : <i>Donkey orchid symptomless virus</i> |
| 2015.012bP | create a new genus in the family <i>Alphaflexiviridae</i> , order <i>Tymovirales</i> |
| 2015.012cP | name the genus <i>Platypuvirus</i> |
| 2015.012dP | designate <i>Donkey orchid symptomless virus</i> as type species of the new genus |
| 2015.013aP.A.v3.Potexvirus_sp [109] | |
| 2015.013aP | create species in the genus <i>Potexvirus</i> , family <i>Alphaflexiviridae</i> , order <i>Tymovirales</i> : <i>Yam virus X</i> |

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 *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 *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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