

## Ratification vote on taxonomic proposals to the International Committee on Taxonomy of Viruses (2012)

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In accordance with the Statutes of the International Committee of Taxonomy of Viruses (ICTV), the final stage in the process of making changes to the Universal Scheme of Virus Classification is the ratification of taxonomic proposals by ICTV members. This can occur either at a plenary meeting of ICTV, held during an International Congress of Virology conference, or by circulation of proposals by mail followed by a ballot. As in recent years, a list of proposals that had been subjected to the full, multi-stage review process was prepared and presented on the ICTVonline web pages by January 2012. This review process involved input from the ICTV study groups and subcommittees, other interested virologists, and the ICTV Executive Committee. These proposals were submitted to and discussed by the ICTV Executive Committee over the past two years and represent changes to virus taxonomy from that recently published in the Ninth Report [1]. The ratification process was performed entirely by email. The proposals were sent electronically via email on January 24 2012 to ICTV Executive Committee Members (18), Life Members (11), ICTV Subcommittee Members (105), and ICTV National Representatives (55). Members were then requested to vote on whether or not to ratify the taxonomic

proposals, within a four-week deadline (voting closed on February 21, 2012).

The following are the taxonomic proposals that were ratified by ICTV members in February 2012. The proposals are organized by subcommittee, and then the filename posted on-line, which contained the specific information about the proposals. Note that by ICTV convention, all approved taxon names are italicized. The details of all these proposals can be obtained at [http://www.ictvonline.org/Official\\_ICTV\\_Proposals.htm](http://www.ictvonline.org/Official_ICTV_Proposals.htm). The latest version of virus taxonomy, incorporating all the changes reported here, is now available on-line at the ICTV web site: <http://ictvonline.org>.

### Fungal Virus Subcommittee

#### 2009.001a-fF.A.v6.Dinodnavirus

- 2009.001a,bF create species named *Heterocapsa circularisquama DNA virus 01* in the new unassigned genus *Dinodnavirus*  
2009.001c,eF create a new unassigned genus  
2009.001dF name the new genus *Dinodnavirus*  
2009.001fF designate *Heterocapsa circularisquama DNA virus 01* as type species in the new genus

#### 2009.002a-fF.A.v6.Bacilladnavirus

- 2009.002a,b,F create species named *Chaetoceros salsugineum DNA virus 01* in the new unassigned genus *Bacilladnavirus*  
2009.002c,eF create a new unassigned genus  
2009.002dF name the new genus *Bacilladnavirus*  
2009.002fF designate *Chaetoceros salsugineum DNA virus 01* as type species in the new genus

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**2010.001a-dF.A.v2.Trichomonasvirus**

- 2010.001aF create species named *Trichomonas vaginalis virus 1*, *Trichomonas vaginalis virus 2*, and *Trichomonas vaginalis virus 3* in the new genus *Trichomonasvirus*, family *Totiviridae*
- 2010.001bF create a genus in the family *Totiviridae*
- 2010.001cF name the new genus *Trichomonasvirus*
- 2010.001dF designate *Trichomonas vaginalis virus 1* as type species in the new genus

**2010.002a-gF.A.v1.Megabirnaviridae**

- 2010.002aF create species named *Rosellinia necatrix megabirnavirus 1* in the new genus *Megabirnavirus* in the new family *Megabirnaviridae*
- 2010.002bF create a genus in the family *Megabirnaviridae*
- 2010.002cF name the new genus *Megabirnavirus*
- 2010.002dF designate *Rosellinia necatrix megabirnavirus 1* as type species in the new genus
- 2010.002eF create an unassigned family
- 2010.002fF name the new family *Megabirnaviridae*
- 2010.002gF assign the new genus *Megabirnavirus* to the new family

**2010.003aF.A.v1.Prasinivirus-sp**

- 2010.003aF create species named *Ostreococcus tauri virus OtV5* in the genus *Prasinivirus*, family *Phycodnaviridae*

**2010.004aF.A.v1.Chlorovirus-Sp**

- 2010.004aF create species named *Acanthocystis turfacea chlorella virus 1* in the genus *Chlorovirus*, family *Phycodnaviridae*

**Invertebrate Virus Subcommittee****2008.001-004,6I.A.v3.Aparavirus**

- 2008.001I create new genus in the family *Dicistroviridae*
- 2008.002I name the new genus *Aparavirus*
- 2008.003I assign 4 existing species (*Acute bee paralysis virus*, *Kashmir bee virus*, *Solenopsis invicta virus-1*, *Taura syndrome virus*) to new genus and create 1 additional species (*Israeli acute paralysis virus*)
- 2008.004I designate *Acute bee paralysis virus* as type species of new genus
- 2008.006I create species Israeli acute paralysis virus in genus *Aparavirus*

**2009.001a-kI.A.v4.Hytrosaviridae**

- 2009.001aI create species named *Glossina hytrovirus* in the proposed genus *Glossinavirus* in the proposed family *Hytrosaviridae*
- 2009.001bI create species named *Musca hytrovirus* in the proposed genus *Muscavirus* in the proposed family *Hytrosaviridae*
- 2009.001cI create a new genus
- 2009.001dI name the genus *Glossinavirus*
- 2009.001eI designate *Glossina hytrovirus* to be type species in the proposed genus *Glossinavirus*
- 2009.001fI create a new genus
- 2009.001gI name the genus *Muscavirus*
- 2009.001hI designate *Musca hytrovirus* to be type species in the proposed genus *Muscavirus*
- 2009.001iI create a new unassigned family
- 2009.001jI name the family *Hytrosaviridae*
- 2009.001kI assign the genera *Glossinavirus* and *Muscavirus* to the new family

**2010.001a-qI.A.v4.Tetraviridae**

- 2010.001aI create a new genus in the family *Permutotetraviridae* (new)
- 2010.001bI name the genus *Alphapermutotetravirus*
- 2010.001cI designate species *Thosea asigna virus* as type species of the new genus
- 2010.001dI create new unassigned family
- 2010.001eI name the new family *Permutotetraviridae*
- 2010.001fI assign genus *Alphapermutotetravirus* to the new family
- 2010.001gI create a new genus in the family *Carmotetraviridae* (new)
- 2010.001hI name the genus *Alphacarmotetravirus*
- 2010.001iI designate species *Providence virus* as type species of the new genus
- 2010.001jI create new unassigned family
- 2010.001kI name the new family *Carmotetraviridae*
- 2010.001lI assign genus *Alphacarmotetravirus* to the new family
- 2010.001mI remove 2 species, *Euprosterina elaeasa virus* and *Thosea asigna virus*, from their current assigned position in the genus *Betatetravirus*, family *Tetraviridae*
- 2010.001nI reassign the species *Euprosterina elaeasa virus* and *Thosea asigna virus* to genus *Alphapermutotetravirus* (new), family *Permutotetraviridae* (new)
- 2010.001oI remove the species *Providence virus* from its current assigned position in the genus *Betatetravirus*, family *Tetraviridae*

- 2010.001pI reassign the species *Providencia virus* to the genus *Alphacarmotetravirus* (new), family *Carmotetraviridae* (new)
- 2010.001qI change the name of family *Tetraviridae* to the *Alphatetraviridae*
- 2011.004aI.A.v1.Aparavirus-Sp**
- 2011.004aI create species named *Mud crab virus* in the genus *Aparavirus*, family *Dicistroviridae*, order *Picornavirales*
- 2011.005aI.A.v1.Iflavirus-Sp**
- 2011.005aI create species named *Slow bee paralysis virus* in the genus *Iflavirus*, family *Iflaviridae*, order *Picornavirales*
- Plant Virus Subcommittee**
- 2009.004a-iP.A.v2.Poacevirus**
- 2009.004a,bP create species named *Triticum mosaic virus* in the new genus *Poacevirus*, family *Potyviridae*
- 2009.004c,eP create a new genus in the family *Potyviridae*
- 2009.004dP name the new genus *Poacevirus*
- 2009.004fP designate *Triticum mosaic virus* as type species in the new genus
- 2009.004gP assign the previously unassigned species *Sugarcane streak mosaic virus* as additional species in the new genus
- 2009.004h,iP remove the species *Sugarcane streak mosaic virus* from “unassigned” in the family *Potyviridae* and reassign to the new genus
- 2009.008a-fP.A.v1.Tepovirus**
- 2009.008a,cP create a genus in the new family *Betaflexiviridae*, order *Tymovirales*
- 2009.008bP name the new genus *Tepovirus*
- 2009.008dP designate *Potato virus T* as type species in the new genus
- 2009.008e,fP remove the species *Potato virus T* from “unassigned” in the family *Betaflexiviridae* and reassign to the new genus
- 2009.012a-fP.A.v6.Labyrnavirus**
- 2009.012a,bP create species named *Aurantiochytrium single-stranded RNA virus 01* in the proposed genus *Labyrnavirus*, to be unassigned in the order *Picornavirales*
- 2009.012c,e,P create an unassigned genus in the order *Picornavirales*
- 2009.012dP name the new genus *Labyrnavirus*
- 2009.012fP designate *Aurantiochytrium single-stranded RNA virus 01* as type species in the new genus
- 2009.015a-kP.A.v6.Bacillarnavirus**
- 2009.015a-f,kP create species named *Rhizosolenia setigera RNA virus 01*, *Chaetoceros tenuissimus RNA virus 01* and *Chaetoceros socialis f. radians RNA virus 01* in the proposed genus *Bacillarnavirus*, unassigned in the order *Picornavirales*
- 2009.015g,I,oP create a new genus to be unassigned in order *Picornavirales*
- 2009.015hP name the new genus *Bacillarnavirus*
- 2009.015jP designate *Rhizosolenia setigera RNA virus 01* as type species in the new genus
- 2009.016a-iP.A.v7.Alvernviridae**
- 2009.016a,bP create species named *Heterocapsa circularisquama RNA virus 01* in the genus *Dinornavirus* (new), family *Alvernviridae* (new, unassigned)
- 2009.016c,e,iP create a genus in the family *Alvernviridae* (new)
- 2009.016eP name the new genus *Dinornavirus*
- 2009.016fP designate *Heterocapsa circularisquama RNA virus 01* as type species in the new genus
- 2010.001aP.A.v2.Ipomovirus-Sp**
- 2010.001aP create species named *Ugandan cassava brown streak virus* in the genus *Ipomovirus* in the family *Potyviridae*
- 2010.002aP.A.v2.Potyvirus-Sp**
- 2010.002aP create species named *Sweet potato virus C* in the genus *Potyvirus* in the family *Potyviridae*
- 2010.003aP.A.v1.Closterovirus-2sp**
- 2010.003aP create 2 species named *Strawberry chlorotic fleck-associated virus* and *Raspberry leaf mottle virus* in the genus *Closterovirus* in the family *Closteroviridae*
- 2010.004aP.A.v1.Cucumovirus-1sp**
- 2010.004aP create species named *Gayfeather mild mottle virus* in the genus *Cucumovirus* in the family *Bromoviridae*

**2010.005aP.A.v1.Ilarvirus-2sp**

2010.005aP create 2 species named *Blackberry chlorotic ringspot virus* and *Strawberry necrotic shock virus* in the genus *Ilarvirus* in the family *Bromoviridae*

*bacilliform virus* to *Dioscorea bacilliform AL virus*, (iii) *Banana streak Mysore virus* to *Banana streak MY virus* and (iv) *Sugarcane bacilliform Mor virus* to *Sugarcane bacilliform MO virus*

**2010.006aP.A.v2.Tymoviridae-sp**

2010.006aP create species *Bombyx mori latent virus* to be unassigned in the family *Tymoviridae*, order *Tymovirales*

**2010.014aP.A.v1.Caulimovirus\_1sp**

2010.014aP create species named *Lamium leaf distortion virus* in the genus *Caulimovirus* in the family *Caulimoviridae*

**2010.007aP.A.v1.Marafivirus-sp\_BIVS**

2010.007aP create species named *Blackberry virus S* in the genus *Marafivirus* in the family *Tymoviridae*, order *Tymovirales*

**2010.015aP.A.v1.Emaravirus\_1sp**

2010.015aP create species named *Fig mosaic virus* in the unassigned genus *Emaravirus*

**2010.008aP.A.v1.Marafivirus-sp\_GSyV-1**

2010.008aP create species named *Grapevine Syrah virus 1* in the genus *Marafivirus* in the family *Tymoviridae*, order *Tymovirales*

**2010.016aP.A.v1.Ilarvirus\_1sp**

2010.016aP create species named *Lilac leaf chlorosis virus* in the genus *Ilarvirus* in the family *Bromoviridae*

**2010.009aP.A.v1.Nanovirus-2sp**

2010.009aP create species named *Faba bean necrotic stunt virus* and *Pea necrotic yellow dwarf virus* in the genus *Nanovirus* in the family *Nanoviridae*

**2010.017a-eP.A.v2.Solendovirus**

2010.017aP create new genus in the family *Caulimoviridae*

2010.017bP name the new genus *Solendovirus*

2010.017cP designate *Tobacco vein clearing virus* as type species in the new genus

2010.017dP remove *Tobacco vein clearing virus* from the genus *Cavemovirus*, family *Caulimoviridae*

2010.017eP reassign *Tobacco vein clearing virus* to the new genus *Solendovirus*

**2010.010aP.A.v1.Pospiviroid-sp**

2010.010aP create species named *Pepper chat fruit viroid* in the genus *Pospiviroid* in the family *Pospiviroidae*

**2010.018aP.A.v1.Soymovirus\_1sp**

2010.018aP create species named *Cestrum yellow leaf curling virus* in the genus *Soymovirus* in the family *Caulimoviridae*

**2010.011aP.A.v1.Remove2unassignedspecies**

2010.011aP remove (abolish) two unassigned species: *Hawaiian rubus leaf curl virus* and *Pigeon pea sterility mosaic virus*

**2010.012aP.A.v1.Tymoviridae-sp**

2010.012aP create species named *Poinsettia mosaic virus* to be unassigned in the family *Tymoviridae*, order *Tymovirales*

**2010.019aP.A.Tymovirus\_1sp**

2010.019aP create species named *Chiltepin yellow mosaic virus* in the genus *Tymovirus* in the family *Tymoviridae*, order *Tymovirales*

**2010.013a,bP.A.v1.Badnavirus\_5sp\_and\_renames**

2010.013aP create 5 species named *Banana streak VN virus*, *Pineapple bacilliform CO virus*, *Pineapple bacilliform ER virus*, *Bougainvillea chlorotic vein banding virus*, and *Dioscorea bacilliform SN virus* in the genus *Badnavirus* in the family *Caulimoviridae*

**2010.020aP.A.v1.Potyvirus\_1sp**

2010.020aP create species named *Sunflower chlorotic mottle virus* in the genus *Potyvirus* in the family *Potyviridae*

2010.013bP Change the names of the following 4 species: (i) *Citrus mosaic virus* to *Citrus yellow mosaic virus*, (ii) *Dioscorea*

**2011.001aP.A.v1.Cavemovirus\_1sp**

2011.001aP create new species named *Sweet potato collusive virus* in the genus *Cavemovirus*, family *Caulimoviridae*

**2011.002aP.A.v1.Solendovirus\_1sp**

2011.002aP create new species named *Sweet potato vein clearing virus* in the genus *Solendovirus* (proposed in <2010.017a-eP.U.v2.Solendovirus>), family *Caulimoviridae*

**2011.003aP.A.v1.Nepovirus\_1sp**

2011.003aP create species *Melon mild mottle virus* in the genus *Nepovirus*, family *Secoviridae*, order *Picornavirales*

**2011.004aP.A.v1.Potyvirus\_1sp**

2011.004aP create species *Yambean mosaic virus* in the genus *Potyvirus*, family *Potyviridae*

**2011.006a,bP.A.v1\_Begomo\_rem,ren\_Sp**

2011.006aP Remove (abolish) 4 species (*Tomato leaf curl Indonesia virus*, *Tomato leaf curl Pakistan virus*, *Squash yellow mild mottle virus* and *Pepper leaf curl Pakistan virus*) from genus *Begomovirus*, family *Geminiviridae*

2011.006bP rename three species in the genus *Begomovirus*: *Corchorus yellow vein Vietnam virus* to *Corchorus yellow vein virus*, *Cucurbit leaf curl virus* to *Cucurbit leaf crumple virus* and *Tomato mosaic leaf curl virus* to *Merremia mosaic virus*

**2011.007aP.A.v1.Carmovirus\_4sp**

2011.007aP create 4 species (*Calibrachoa mottle virus*, *Honeysuckle ringspot virus*, *Nootka lupine vein clearing virus* and *Soybean yellow mottle mosaic virus*) in the genus *Carmovirus*, family *Tombusviridae*

**2011.008aP.A.v1.Panicovirus\_1sp**

2011.008aP create species *Cocksfoot mild mosaic virus* in the genus *Panicovirus*, family *Tombusviridae*

**2011.011aP.A.v1.Sobemovirus\_1sp**

2011.011aP create species named *Imperata yellow mottle virus* in the genus *Sobemovirus*

**2011.012aP.A.v1Furovirus\_1sp**

2011.012aP create species named *Japanese soil-borne wheat mosaic virus* in the genus *Furovirus* in the family *Virgaviridae*

**2011.013aP.A.v1.Badnavirus\_2sp**

2011.013aP create 2 species, *Sweet potato pakakuy virus* and *Grapevine vein clearing virus*, in the genus *Badnavirus* in the family *Caulimoviridae*

**2011.014aP.A.v1.Marafivirus\_1sp**

2011.014aP create species named *Olive latent virus 3* in the genus *Marafivirus* in the family *Tymoviridae*, order *Tymovirales*

**Prokaryote virus subcommittee****2009.001a-gB.A.v2.PB1likevirus**

2009.001a,b,gB create species named *Pseudomonas phage PB1*, *Pseudomonas phage SN*, *Pseudomonas phage 14-1*, *Pseudomonas phage LMA2*, *Pseudomonas phage LBL3*, *Pseudomonas phage F8*, and *Burkholderia phage BcepF1* in the new genus *PB1likevirus*, family *Myoviridae*, order *Caudovirales*

2009.001c,eB create a new genus in the family *Myoviridae*

2009.001dB name the new genus *PB1likevirus*

2009.001fB designate *Pseudomonas phage PB1* as type species in the new genus

**2009.002a-gB.A.v2.Bcep781likevirus**

2009.002a,b,gB create species named *Burkholderia phage Bcep781*, *Burkholderia phage Bcep43*, *Burkholderia phage Bcep1*, *Burkholderia phage BcepNY3*, and *Xanthomonas phage OP2* in the new genus *Bcep781likevirus*, family *Myoviridae*

2009.002c,eB create a new genus in the family *Myoviridae*

2009.002dB name the new genus *Bcep781likevirus*

2009.002fB designate *Burkholderia phage Bcep781* as type species in the new genus

**2009.003a-gB.A.v2.Bcepmulikevirus**

2009.003a,b,gB create species named *Burkholderia phage BcepMu* and *Burkholderia phage phiE255* in the new genus *Bcepmulikevirus*, family *Myoviridae*

2009.003c,eB create a new genus in the family *Myoviridae*

- 2009.003dB name the new genus *Bcepmulikevirus*  
 2009.003fB designate *Burkholderia phage BcepMu* as type species in the new genus
- 2009.004a-gB.A.v2.Hap1likevirus**
- 2009.004a,b,gB create species named *Halomonas phage phiHAP-1* and *Vibrio phage VP882* in the new genus *Hap1likevirus*, family *Myoviridae*  
 2009.004c,eB create a new genus in the family *Myoviridae*  
 2009.004dB name the new genus *Hap1likevirus*  
 2009.004fB designate *Halomonas phage phiHAP-1* as type species in the new genus
- 2009.005a-gB.A.v2.PhiCD119likevirus**
- 2009.005a,b,gB create species named *Clostridium phage phiCD119*, *Clostridium phage phiC2* and *Clostridium phage phiCD27* in the new genus *PhiCD119likevirus*, family *Myoviridae*  
 2009.005c,eB create a new genus in the family *Myoviridae*  
 2009.005dB name the new genus *PhiCD119likevirus*  
 2009.005fB designate *Clostridium phage phiCD119* as type species in the new genus
- 2009.006a-gB.A.v2.FelixO1likevirus**
- 2009.006a,b,gB create species named *Salmonella phage FelixO1*, *Erwinia phage phiEa21-4* and *Escherichia phage wV8* in the new genus *FelixO1likevirus*, family *Myoviridae*  
 2009.006c,eB create a new genus in the family *Myoviridae*  
 2009.006dB name the new genus *FelixO1likevirus*  
 2009.006fB designate *Salmonella phage FelixO1* as type species in the new genus
- 2009.007a-rB.A.v3.Tevenvirinae**
- 2009.007a,b,mB create 10 new species (names: *Escherichia phage JS98*, *Escherichia phage RB14*, *Escherichia phage RB16*, *Escherichia phage RB32*, *Escherichia phage RB43*, *Escherichia phage RB49*, *Escherichia phage RB69*, *Escherichia phage phi1*, *Aeromonas phage 31*, *Aeromonas phage 25*) in the genus “T4-like viruses” (proposed name *T4likevirus*), subfamily *Tevenvirinae* (new), family *Myoviridae*, order *Caudovirales*  
 2009.007c,dB create a new species named *Vibrio phage KVP40* in the genus *Schizot4likevirus* (new), subfamily *Tevenvirinae* (new), family *Myoviridae*, order *Caudovirales*  
 2009.007e,g,mB create a new genus in the subfamily *Tevenvirinae* (new), family *Myoviridae*, order *Caudovirales*  
 2009.007fB name the new genus *Schizot4likevirus*  
 2009.007hB designate *Vibrio phage KVP40* as type species in the new genus  
 2009.007iB assign *Vibrio phage nt-1* as an additional species in the new genus  
 2009.007j,lB create a new subfamily in family *Myoviridae*, order *Caudovirales*  
 2009.007kB name the new subfamily *Tevenvirinae*  
 2009.007nB assign *Acinetobacter phage 133*, *Aeromonas phage Aeh1* and *Aeromonas phage 65* as unassigned species in the new subfamily *Tevenvirinae*  
 2009.007o,pB move the species *Vibrio phage nt-1* from the genus “T4-like viruses” to the genus *Schizot4likevirus* (new)  
 2009.007q,rB remove the species *Acinetobacter phage 133*, *Aeromonas phage Aeh1* and *Aeromonas phage 65* from the genus “T4-like viruses” and designate as unassigned in the new subfamily *Tevenvirinae*
- 2009.009a-pB.A.v3.Spounavirinae**
- 2009.009a,b,mB create 5 new species (names: *Staphylococcus phage Twort*, *Staphylococcus phage K*, *Staphylococcus phage G1*, *Listeria phage P100*, *Listeria phage A511*) in the genus *Twortlikevirus*, subfamily *Spounavirinae* (new), family *Myoviridae*, order *Caudovirales*  
 2009.009c,dB create 2 new species named *Enterococcus phage phiEC24C* and *Lactobacillus phage LP65* to be unassigned in the subfamily *Spounavirinae* (new), family *Myoviridae*, order *Caudovirales*  
 2009.009e,g,mB create a new genus in the subfamily *Spounavirinae* (new), family *Myoviridae*, order *Caudovirales*  
 2009.009fB name the new genus *Twortlikevirus*  
 2009.009hB designate *Staphylococcus phage Twort* as type species in the new genus

- 2009.009iB assign *Staphylococcus phage K*, *Staphylococcus phage G1*, *Listeria phage P100* and *Listeria phage A511* as additional species in the new genus
- 2009.009j,IB create a new subfamily in family *Myoviridae*, order *Caudovirales*
- 2009.009kB name the new subfamily *Spounavirinae*
- 2009.009nB assign *Enterococcus phage phiEC24C* and *Lactobacillus phage LP65* as unassigned species in the new subfamily *Spounavirinae*
- 2009.009o,pB move genus “SPO1-like viruses” (proposed name *SPO1likevirus*), currently unassigned in the family *Myoviridae*, into the new subfamily *Spounavirinae*
- 2009.012a-qV.A.v2.Peduovirinae**
- 2009.012a,b,mB create 12 new species (names: *Enterobacteria phage Wphi*, *Yersinia phage L-413C*, *Enterobacteria phage 186*, *Enterobacteria phage PsP3*, *Salmonella Fels-2*, *Salmonella Sop-Ephi*, *Burkholderia phage phiE202*, *Mannheimia phage phiMhaA1-PHL101*, *Pseudomonas phage phiCTX*, *Burkholderia phage phi52237*, *Ralstonia phage RSA1*, *Burkholderia phage phiE12-2*) in the genus “P2-like viruses” (proposed name *P2likevirus*), subfamily *Peduovirinae* (new), family *Myoviridae*, order *Caudovirales*
- 2009.012c,d,iB create 6 new species (names: *Haemophilus phage HP1*, *Haemophilus phage HP2*, *Pasteurella phage F108*, *Vibrio phage K139*, *Vibrio phage Kappa*, *Aeromonas phage phiO18P*) in the genus *Hp1likevirus* (new), subfamily *Peduovirinae* (new), family *Myoviridae*, order *Caudovirales*
- 2009.012e,g,mB create a new genus in the subfamily *Peduovirinae* (new), family *Myoviridae*, order *Caudovirales*
- 2009.012fB name the new genus *Hp1likevirus*
- 2009.012hB designate *Haemophilus phage HP1* as type species in the new genus
- 2009.012j,IB create a new subfamily in the family *Myoviridae*, order *Caudovirales*
- 2009.012kB name the new subfamily *Peduovirinae*
- 2009.012n,oB move species *Haemophilus phage HP1*, currently assigned to the genus “P2-like viruses” (proposed name *P2likevirus*), family *Myoviridae*, into the new genus *HP1likevirus*
- 2009.012p,qB move genus “P2-like viruses” (proposed name *P2likevirus*), currently unassigned in the family *Myoviridae*, into the new subfamily *Peduovirinae*
- 2010.002a-gB.A.v2.Clavaviridae**
- 2010.002aB create species named *Aeropyrum pernix bacilliform virus 1* in the new genus *Clavavirus*, family *Clavaviridae*
- 2010.002bB create a genus in the family *Clavaviridae*
- 2010.002cB name the new genus *Clavavirus*
- 2010.002dB designate *Aeropyrum pernix bacilliform virus 1* as type species in the new genus
- 2010.002eB create an unassigned family
- 2010.002fB name the new family *Clavaviridae*
- 2010.002gB assign the new genus *Clavavirus* to the new family
- 2011.002aB.A.v2.Remove\_EnterophageEra103**
- 2011.002aB Remove (abolish) the species *Enterobacteria phage Era103* from the genus “SP6-like viruses” (proposed new genus name *Sp6likevirus*)
- 2011.003aB.A.v2.Rename\_phiKMV-like\_species**
- 2011.003aB in the genus “PhiKMV-like viruses” (proposed name *Phikmvlikevirus*) change the name of species *Enterobacteria phage phiKMV* to *Pseudomonas phage phiKMV* and *Enterobacteria phage LKA1* to *Pseudomonas phage LKA1*
- 2011.004abB.A.v2.Correct\_AHJD-like\_errors**
- 2011.004aB Remove (abolish) species *44AHJD* from genus “AHJD-like viruses” (proposed name *Ahjdlikevirus*), subfamily *Picovirinae*, family *Podoviridae*
- 2011.004bB change the name of species *Staphylococcus phage AHJD* to *Staphylococcus phage 44AHJD*
- 2011.005abB.A.v2.Correct\_BPP-1-like\_errors**
- 2011.005aB remove (abolish) species *Salmonella phage BPP-1* from genus “Bpp-1-like viruses” (proposed name *Bpp1likevirus*), family *Podoviridae*
- 2011.005aB designate *Bordetella phage BPP-1* as the type species of the genus “BPP-1-like viruses”

<b>2011.006aB.A.v2.Ren_Myobacteria_phage_D29_sp</b>	2010.001cV	create species named <i>Deltapapillomavirus</i> 5 in the genus <i>Deltapapillomavirus</i> , family <i>Papillomaviridae</i>
2011.006aB in the genus “L5-like viruses” (proposed name <i>L5likevirus</i> ) change the name of species <i>Mycobacteria phage D29</i> to <i>Mycobacterium phage D29</i>	2010.001dV	create 2 species named <i>Lambdapapillomavirus</i> 3 and <i>Lambdapapillomavirus</i> 4 in the genus <i>Lambdapapillomavirus</i> , family <i>Papillomaviridae</i>
<b>2011.007aB.A.v2.Ren_Enterobacteria_phage_N4_sp</b>	2010.001eV	create species named <i>Pipapillomavirus</i> 2 in the genus <i>Pipapillomavirus</i> , family <i>Papillomaviridae</i>
2011.007aB in the genus “N4-like viruses”, family <i>Podoviridae</i> , change the name of species <i>Enterobacteria phage N4</i> to <i>Escherichia phage N4</i>	2010.001fV	create species named <i>Rhopapillomavirus</i> 1 in the proposed genus <i>Rhopapillomavirus</i> , family <i>Papillomaviridae</i>
<b>Vertebrate virus subcommittee</b>	2010.001gV	create species named <i>Sigmapapillomavirus</i> 1 in the proposed genus <i>Sigmapapillomavirus</i> , family <i>Papillomaviridae</i>
<b>2009.005a-fV.A.v2.Aquaparamyxovirus</b>		
2009.005a,bV create species, named <i>Atlantic salmon paramyxovirus</i> , in the new genus <i>Aquaparamyxovirus</i> , subfamily <i>Paramyxovirinae</i> , family <i>Paramyxoviridae</i> , order <i>Mononegavirales</i>	2010.001hV	create species named <i>Taupapillomavirus</i> 1 in the proposed genus <i>Taupapillomavirus</i> , family <i>Papillomaviridae</i>
2009.005c,eV create a new genus in the subfamily <i>Paramyxovirinae</i>	2010.001iV	create 2 species named <i>Upsilonpapillomavirus</i> 1 and <i>Upsilonpapillomavirus</i> 2 in the proposed genus <i>Upsilonpapillomavirus</i> , family <i>Papillomaviridae</i>
2009.005dV name the new genus <i>Aquaparamyxovirus</i>	2010.001jV	create species named <i>Phipapillomavirus</i> 1 in the proposed genus <i>Phipapillomavirus</i> , family <i>Papillomaviridae</i>
2009.005fV designate <i>Atlantic salmon paramyxovirus</i> as the type species of the new genus	2010.001kV	create 2 species named <i>Chipapillomavirus</i> 1 and <i>Chipapillomavirus</i> 2 in the proposed genus <i>Chipapillomavirus</i> , family <i>Papillomaviridae</i>
<b>2009.007a-fV.A.v2.Ferlavirus</b>		
2009.007a,bV create species, named <i>Fer-de-Lance paramyxovirus</i> , in the new genus <i>Ferlavirus</i> , subfamily <i>Paramyxovirinae</i> , family <i>Paramyxoviridae</i> , order <i>Mononegavirales</i>	2010.001lV	create species named <i>Psipapillomavirus</i> 1 in the proposed genus <i>Psipapillomavirus</i> , family <i>Papillomaviridae</i>
2009.007c,eV create a new genus in the subfamily <i>Paramyxovirinae</i>	2010.001mV	create species named <i>Omegapapillomavirus</i> 1 in the proposed genus <i>Omegapapillomavirus</i> , family <i>Papillomaviridae</i>
2009.007dV name the new genus <i>Ferlavirus</i>	2010.001nV	create species named <i>Dyodeltapapillomavirus</i> 1 in the proposed genus <i>Dyodeltapapillomavirus</i> , family <i>Papillomaviridae</i>
2009.007fV designate <i>Fer-de-Lance paramyxovirus</i> as the type species of the new genus		
<b>2010.001a-kkkV.A.v2.Papillomaviridae</b>		
2010.001aV create species named <i>Betapapillomavirus</i> 6 in the genus <i>Betapapillomavirus</i> , family <i>Papillomaviridae</i>	2010.001oV	create species named <i>Dyoepsilonpapillomavirus</i> 1 in the proposed genus <i>Dyoepsilonpapillomavirus</i> , family <i>Papillomaviridae</i>
2010.001bV create 5 species ( <i>Gammapapillomavirus</i> 6, <i>Gammapapillomavirus</i> 7, <i>Gammapapillomavirus</i> 8, <i>Gammapapillomavirus</i> 9 and <i>Gammapapillomavirus</i> 10) in the genus <i>Gammapapillomavirus</i> , family <i>Papillomaviridae</i>	2010.001pV	create species named <i>Dyozetapapillomavirus</i> 1 in the proposed genus <i>Dyozetapapillomavirus</i> , family <i>Papillomaviridae</i>



2010.001qV	create species named <i>Dyoetapapillomavirus 1</i> in the proposed genus <i>Dyoetapapillomavirus</i> , family <i>Papillomaviridae</i>	2010.001ssV	name the new genus <i>Dyodeltapapillomavirus</i>
2010.001rV	create species named <i>Dyothetapapillomavirus 1</i> in the proposed genus <i>Dyothetapapillomavirus</i> , family <i>Papillomaviridae</i>	2010.001ttV	designate <i>Dyodeltapapillomavirus 1</i> as type species in the new genus
2010.001sV	create species named <i>Dyoiotapapillomavirus 1</i> in the proposed genus <i>Dyoiotapapillomavirus</i> , family <i>Papillomaviridae</i>	2010.001uuV	create new genus within the family <i>Papillomaviridae</i>
2010.001tV	create new genus within the family <i>Papillomaviridae</i>	2010.001vvV	name the new genus <i>Dyoepsilonpapillomavirus</i>
2010.001uV	name the new genus <i>Rhopapillomavirus</i>	2010.001wwV	designate <i>Dyoepsilonpapillomavirus 1</i> as type species in the new genus
2010.001vV	designate <i>Rhopapillomavirus 1</i> as type species in the new genus	2010.001xxV	create new genus within the family <i>Papillomaviridae</i>
2010.001wV	create new genus within the family <i>Papillomaviridae</i>	2010.001yyV	name the new genus <i>Dyozetapapillomavirus</i>
2010.001xV	name the new genus <i>Sigmapapillomavirus</i>	2010.001zzV	designate <i>Dyozetapapillomavirus 1</i> as type species in the new genus
2010.001yV	designate <i>Sigmapapillomavirus 1</i> as type species in the new genus	2010.001aaaV	create new genus within the family <i>Papillomaviridae</i>
2010.001zV	create new genus within the family <i>Papillomaviridae</i>	2010.001bbbV	name the new genus <i>Dyoetapapillomavirus</i>
2010.001aaV	name the new genus <i>Taupapillomavirus</i>	2010.001cccV	designate <i>Dyoetapapillomavirus 1</i> as type species in the new genus
2010.001bbV	designate <i>Taupapillomavirus 1</i> as type species in the new genus	2010.001dddV	create new genus within the family <i>Papillomaviridae</i>
2010.001ccV	create new genus within the family <i>Papillomaviridae</i>	2010.001eeeV	name the new genus <i>Dyothetapapillomavirus</i>
2010.001ddV	name the new genus <i>Upsilonpapillomavirus</i>	2010.001fffV	designate <i>Dyothetapapillomavirus 1</i> as type species in the new genus
2010.001eeV	designate <i>Upsilonpapillomavirus 1</i> as type species in the new genus	2010.001gggV	create new genus within the family <i>Papillomaviridae</i>
2010.001ffV	create new genus within the family <i>Papillomaviridae</i>	2010.001hhhV	name the new genus <i>Dyoiotapapillomavirus</i>
2010.001ggV	name the new genus <i>Phipapillomavirus</i>	2010.001iiiV	designate <i>Dyoiotapapillomavirus 1</i> as type species in the new genus
2010.00hhV	designate <i>Phipapillomavirus 1</i> as type species in the new genus	2010.001jjjV	Remove (abolish) species <i>Human papillomavirus 71</i>
2010.001iiV	create new genus within the family <i>Papillomaviridae</i>	2010.001kkkV	rename 43 species in the family <i>Papillomaviridae</i> ( <i>Human papillomavirus 32</i> to <i>Alphapapillomavirus 1</i> ; <i>Human papillomavirus 10</i> to <i>Alphapapillomavirus 2</i> ; <i>Human papillomavirus 61</i> to <i>Alphapapillomavirus 3</i> ; <i>Human papillomavirus 2</i> to <i>Alphapapillomavirus 4</i> ; <i>Human papillomavirus 26</i> to <i>Alphapapillomavirus 5</i> ; <i>Human papillomavirus 53</i> to <i>Alphapapillomavirus 6</i> ; <i>Human papillomavirus 18</i> to <i>Alphapapillomavirus 7</i> ; <i>Human papillomavirus 7</i> to <i>Alphapapillomavirus 8</i> ; <i>Human papillomavirus 16</i> to <i>Alphapapillomavirus 9</i> ; <i>Human papillomavirus 6</i> to <i>Alphapapillomavirus 10</i> ; <i>Human</i>
2010.001jjV	name the new genus <i>Chipapillomavirus</i>		
2010.001kkV	designate <i>Chipapillomavirus 1</i> as type species in the new genus		
2010.001llV	create new genus within the family <i>Papillomaviridae</i>		
2010.001mmV	name the new genus <i>Psipapillomavirus</i>		
2010.001nnV	designate <i>Psipapillomavirus 1</i> as type species in the new genus		
2010.001ooV	create new genus within the family <i>Papillomaviridae</i>		
2010.001ppV	name the new genus <i>Omegapapillomavirus</i>		
2010.001qqV	designate <i>Omegapapillomavirus 1</i> as type species in the new genus		
2010.001rrV	create new genus within the family <i>Papillomaviridae</i>		

*papillomavirus 34 to Alphapapillomavirus 11; Rhesus monkey papillomavirus 1 to Alphapapillomavirus 12; Human papillomavirus 54 to Alphapapillomavirus 13; Human papillomavirus cand90 to Alphapapillomavirus 14; Human papillomavirus 5 to Betapapillomavirus 1; Human papillomavirus 9 to Betapapillomavirus 2; Human papillomavirus 49 to Betapapillomavirus 3; Human papillomavirus cand92 to Betapapillomavirus 4; Human papillomavirus cand96 to Betapapillomavirus 5; Human papillomavirus 4 to Gammapapillomavirus 1; Human papillomavirus 48 to Gammapapillomavirus 2; Human papillomavirus 50 to Gammapapillomavirus 3; Human papillomavirus 60 to Gammapapillomavirus 4; Human papillomavirus 88 to Gammapapillomavirus 5; European elk papillomavirus to Deltapapillomavirus 1; Deer papillomavirus to Deltapapillomavirus 2; Ovine papillomavirus 1 to Deltapapillomavirus 3; Bovine papillomavirus 1 to Deltapapillomavirus 4; Bovine papillomavirus 5 to Epsilon papillomavirus 1; Equine papillomavirus 1 to Zetapapillomavirus 1; Fringilla coelebs papillomavirus to Etapapillomavirus 1; Psittacus erithacus timneh papillomavirus to thetapapillomavirus 1; Mastomys natalensis papillomavirus to Iotapapillomavirus 1; Rabbit oral papillomavirus to Kappapapillomavirus 1; Cottontail rabbit papillomavirus to Kappapapillomavirus 2; Feline papillomavirus to Lambdapapillomavirus 1; Canine oral papillomavirus to Lambdapapillomavirus 2; Human papillomavirus 1 to Mupapillomavirus 1; Human papillomavirus 63 to Mupapillomavirus 2; Human papillomavirus 41 to Nupapillomavirus 1; Bovine papillomavirus 3 to Xipapillomavirus 1; Phocoena spinipinnis papillomavirus to Omikronpapillomavirus 1; Hamster oral papillomavirus to Pipapillomavirus 1)*

#### 2010.003aV.A.v2.Nucleorhabdovirus-sp

2010.003aV create species named *Maize Iranian mosaic virus* in the genus *Nucleorhabdovirus*, family *Rhabdoviridae*

#### 2010.004a-dV.A.v2.Lambdatorquevirus

2010.004aV create species named *Torque teno zalophus virus 1* in the proposed genus *Lambdatorquevirus*, family *Anelloviridae*

2010.004bV create new genus within the family *Anelloviridae*

2010.004cV name the new genus *Lambdatorquevirus*

2010.004dV designate *Torque teno zalophus virus 1* as type species in the new genus

#### 2010.005a-dV.A.v3.Kappatorquevirus

2010.005aV create species named *Torque teno sus virus k2* in the proposed genus *Kappatorquevirus*, family *Anelloviridae*

2010.005bV create new genus within the family *Anelloviridae*

2010.005cV name the new genus *Kappatorquevirus*

2010.005dV designate *Torque teno sus virus k2* as type species in the new genus

#### 2010.006aV.A.v2.Etatorquevirus-sp

2010.006aV create species named *Torque teno felis virus 2* in the genus *Etatorquevirus*, family *Anelloviridae*

#### 2010.007aV.A.v2.Betatorquevirus-3sp

2010.007aV create 3 species named *Torque teno mini virus 10*, *Torque teno mini virus 11* and *Torque teno mini virus 12* in the genus *Betatorquevirus*, family *Anelloviridae*

#### 2010.008aV.A.v2.Gammatorquevirus-13sp

2010.008aV create 13 species (*Torque teno midi virus 3*, *Torque teno midi virus 4*, *Torque teno midi virus 5*, *Torque teno midi virus 6*, *Torque teno midi virus 7*, *Torque teno midi virus 8*, *Torque teno midi virus 9*, *Torque teno midi virus 10*, *Torque teno midi virus 11*, *Torque teno midi virus 12*, *Torque teno midi virus 13*, *Torque teno midi virus 14*, *Torque teno midi virus 15*) in the genus *Gammatorquevirus*, family *Anelloviridae*

#### 2010.009aV.A.v2.Rhabdoviridae-sp

2010.009aV create species named *Moussa virus* unassigned in the family *Rhabdoviridae*, order *Mononegavirales*

#### 2010.010aV.A.v1.Ebolavirus-sp

2010.010aV create species named *Bundibugyo ebolavirus* in the genus *Ebolavirus*, family *Filoviridae*, order *Mononegavirales*

**2010.010bV.A.v3.Ebolavirus-ren-Sp**

2010.010bV replace the species name *Lake Victoria marburgvirus* by *Marburg marburgvirus* in the genus *Marburgvirus*

**2010.011aV.A.v1.Mastadenovirus-sp**

2010.011aV create species named *Bat adenovirus A* in the genus *Mastadenovirus*, family *Adenoviridae*

**2010.012aV.A.v1.Mastadenovirus-ren5sp**

2010.012aV rename 5 species in the genus *Mastadenovirus*, family *Adenoviridae*, by adding a letter designation

**2010.014aV.A.v1.Mardivirus-sp**

2010.014aV create species named *Anatid herpesvirus 1* in the genus *Mardivirus*, subfamily *Alphaherpesvirinae*, family *Herpesviridae*, order *Herpesvirales*

**2010.015aV.A.v1.Cyprinivirus-sp**

2010.015aV create species *Anguillid herpesvirus 1* in the genus *Cyprinivirus*, family *Alloherpesviridae*, order *Herpesvirales*

**2010.016a-eV.A.v2.Scutavirus**

2010.016aV create new genus within the subfamily *Alphaherpesvirinae*, family *Herpesviridae*, order *Herpesvirales*

2010.016bV name the new genus *Scutavirus*

2010.016cV designate *Chelonid herpesvirus 5* as type species in the new genus

2010.016dV remove species *Chelonid herpesvirus 5* from its unassigned position in the subfamily *Alphaherpesvirinae*, family *Herpesviridae*

2010.016eV reassign the above species to the new genus

**2010.017a-cV.A.v3.Avastrovirus**

2010.017aV create 3 species named *Avastrovirus 1*, *Avastrovirus 2* and *Avastrovirus 3* in the genus *Avastrovirus*, family *Astroviridae*

2010.017bV remove (abolish) the species *Chicken astrovirus*, *Duck astrovirus* and *Turkey astrovirus* from the genus *Avastrovirus*, family *Astroviridae*

2011.017cV designate *Avastrovirus 1* as the type species of the genus

**2010.018a-cV.A.v4.Mamastrovirus**

2010.018aV create 14 species (*Mamastrovirus 4*, *Mamastrovirus 5*, *Mamastrovirus 6*, *Mamastrovirus 7*, *Mamastrovirus 8*, *Mamastrovirus 9*,

*Mamastrovirus 11*, *Mamastrovirus 12*, *Mamastrovirus 14*, *Mamastrovirus 15*, *Mamastrovirus 16*, *Mamastrovirus 17*, *Mamastrovirus 18*, *Mamastrovirus 19*) in the genus *Mamastrovirus*, family *Astroviridae*

2010.018bV remove (abolish) the species *Bovine astrovirus* from the genus *Mamastrovirus*, family *Astroviridae*

2010.018cV change the names of 5 species in the genus *Mamastrovirus* as follows: *Feline astrovirus* to *Mamastrovirus 2*, *Human astrovirus* to *Mamastrovirus 1*, *Mink astrovirus* to *Mamastrovirus 10*, *Ovine astrovirus* to *Mamastrovirus 13*, *Porcine astrovirus* to *Mamastrovirus 3*

**2010.019a-eV.A.v1.Crocodylidpoxvirus**

2010.019aV create new species named *Skunkpox virus* within the genus *Orthopoxvirus*, subfamily *Chordopoxvirinae*, family *Poxviridae*

2010.019bV create new species named *Nile crocodilepox virus* within the proposed genus *Crocodylidpoxvirus*, subfamily *Chordopoxvirinae*, family *Poxviridae*

2010.019cV create new genus within the subfamily *Chordopoxvirinae*, family *Poxviridae*

2010.019dV name the new genus *Crocodylidpoxvirus*

2010.019eV designate *Nile crocodilepox virus* as type species in the new genus

**2010.020abV.A.v1.Poxviridae-2sp\_ren**

2010.020aV replace the species name *Squirrel poxvirus* by *Squirrelpox virus* in the subfamily *Chordopoxvirinae*, family *Poxviridae*

2010.020bV replace the species name *Deerpox virus W-848-83* by *Mule deerpox virus* in the genus *Cervidpoxvirus*, family *Poxviridae*

**2010.021a-gV.A.v1.Bidnaviridae**

2010.021aV create species named *Bombyx mori bidensovirus* in the proposed genus *Bidensovirus*, in the proposed family *Bidnaviridae*

2010.021bV create new genus within the family *Bidnaviridae*

2010.021cV name the new genus *Bidensovirus*

2010.021dV designate *Bombyx mori bidensovirus* as type species in the new genus

2010.021eV create new unassigned family

2010.021fV name the family *Bidnaviridae*

2010.021gV assign the genus *Bidensovirus* to the new family

**2010.023a-dV.U.v2.Deltacoronavirus**

- 2010.023aV create 3 new species named *Bulbul coronavirus HKU11*, *Thrush coronavirus HKU12* and *Munia coronavirus HKU13* in the proposed genus *Deltacoronavirus*, in the subfamily *Coronavirinae*, family *Coronaviridae*
- 2010.023bV create a new genus in the subfamily *Coronavirinae*, family *Coronaviridae*
- 2010.023cV name the new genus *Deltacoronavirus*
- 2010.023dV designate *Bulbul coronavirus HKU11* as type species in the new genus

**2010.024a,bV.A.v1.Phlebovirus\_2sp-ren**

- 2010.024aV change the name *Chandiru virus*, a species in the genus *Phlebovirus*, to *Candiru virus*
- 2010.024bV change the name *Salehebad virus*, a species in the genus *Phlebovirus*, to *Salehabad virus*

**2011.002aV.A.v2.Anelloviridae\_2Sp-ren**

- 2011.002aV change the names of the following two species: *Torque teno sus virus 1* to *Torque teno sus virus 1a* (genus *Iotatorquevirus*) and *Torque teno sus virus 2* to *Torque teno sus virus 1b* (genus *Iotatorquevirus*)

**2011.003aV.A.v2.Arenavirus-Sp**

- 2011.003aV create species named *Lujo virus* in genus *Arenavirus*, family *Arenaviridae*

**2011.004aV.A.v3.Rhadinovirus-Sp**

- 2011.004aV create species *Murid herpesvirus 7* in the genus *Rhadinovirus*, subfamily *Gammaherpesvirinae*, family *Herpesviridae*, order *Herpesvirales*

**2011.005a-cV.A.v2.Roseolovirus-2sp,rensp**

- 2011.005aV create species *Human herpesvirus 6A* and *Human herpesvirus 6B* in the genus *Roseolovirus*, subfamily *Betaherpesvirinae*, family *Herpesviridae*, order *Herpesvirales*
- 2011.005bV Remove (abolish) species *Human herpesvirus 6* from genus *Roseolovirus*
- 2011.005cV designate *Human herpesvirus 6A* as the type species of the genus

**2011.006aV.A.v1.Lyssavirus-Sp**

- 2011.006aV create species named *Shimoni bat virus* in the genus *Lyssavirus*

**2011.013aV.A.v3.Rhadinovirus-Sp**

- 2011.013aV create species *Cricetid herpesvirus 2* in the genus *Rhadinovirus*, subfamily *Gammaher-*

*pesvirinae*, family *Herpesviridae*, order *Herpesvirales*

**2011.014aV.A.v2.Muromegalovirus-Sp**

- 2011.014aV Create species *Murid herpesvirus 8* in the genus *Muromegalovirus*, subfamily *Betaherpesvirinae*, family *Herpesviridae*, order *Herpesvirales*

**2011.016aV.A.v2.Aphthovirus-Sp**

- 2011.016aV create species *Bovine rhinitis A virus* within the genus *Aphthovirus*, family *Picornaviridae*

**2011.023aV.A.v1.Aviadenovirus-Sp**

- 2011.023aV Create species *Turkey adenovirus B* in the genus *Aviadenovirus*, family *Adenoviridae*

**2011.024aV.A.v1.Mastadenovirus-2Sp**

- 2011.024aV Create two species, *Bat adenovirus B* and *Murine adenovirus B*, in the genus *Mastadenovirus*, family *Adenoviridae*

**2011.025aV.A.v1.Siadenovirus-Sp**

- 2011.025aV Create species *Great tit adenovirus A* in the genus *Siadenovirus*, family *Adenoviridae*

**Conclusion**

All of the above taxonomic proposals were unanimously (or nearly unanimously) approved by the membership who voted. A number of suggestions were received during the voting process. These included requests to rationalize species demarcation criteria across virus families, to pay continued attention to clarity in naming new genera and to seek multiple species demarcation criteria so that new species are not created solely on the basis of a measure of nucleotide sequence identity. The ICTV continues to discuss these issues and welcomes comments on all taxonomic proposals as they are posted on the ICTV web site. The return rate of votes was approximately 48%. These taxonomic changes are now part of the official ICTV taxonomy and supplement the taxonomy presented in the Ninth ICTV Report published in 2011 [1]. An up-to-date list of all approved taxa can be found on the ICTV online web site.

**References**

1. King AMQ, Adams MJ, Carstens EB, Lefkowitz EJ (2011) Virus taxonomy. Ninth report of the international committee on taxonomy of viruses. Elsevier Academic Press, London, San Diego