



This form should be used for all taxonomic proposals. Please complete all those modules that are applicable (and then delete the unwanted sections).

Code(s) assigned:	2007.075a- xxV	(to be completed by ICTV officers)			
Short title: Create the family Anelloviridae containing 9 new genera and 47 new species (e.g. 6 new species in the genus <i>Zetavirus</i> ; re-classification of the family <i>Zetaviridae</i> etc.)					
Modules attached (please check all that apply):	1 <input type="checkbox"/>	2 <input checked="" type="checkbox"/>	3 <input type="checkbox"/>	4 <input checked="" type="checkbox"/>	5 <input checked="" type="checkbox"/>
	6 <input checked="" type="checkbox"/>	7 <input type="checkbox"/>			

Author(s) with e-mail address(es) of the proposer:

Philippe Biagini on behalf of the Circoviridae-Anellovirus Study Group (pbiagini-ets-ap@gulliver.fr)

ICTV-EC or Study Group comments and response of the proposer:

--



MODULE 2: **NEW FAMILY**

Code	2007.075aV	(assigned by ICTV officers)
To create a new family assigned to the order: <i>Unassigned</i>		

Code	2007.075bV	(assigned by ICTV officers)
To name the new family: <i>Anelloviridae</i>		

Code	2007.075cV	(assigned by ICTV officers)
To assign the following genera to the new family:		
<i>Alphatorquevirus</i>		
<i>Betatorquevirus</i>		
<i>Gammatorquevirus</i>		
<i>Deltatorquevirus</i>		
<i>Epsilontorquevirus</i>		
<i>Zetatorquevirus</i>		
<i>Etatorquevirus</i>		
<i>Thetatorquevirus</i>		
<i>Iotatorquevirus</i>		
[All new]		

Code		(assigned by ICTV officers)
To assign the following species to be unassigned in the new family (i.e. within the family but not assigned to any genus):		
.		

Argument to justify the creation of a new family:

The floating genus *Anellovirus* potentially comprises a large and growing number of viruses that share a similar genome organization but are extremely variable in sequence. This prompts the raising of this taxon to that of family and the creation of new genera and species.

Origin of the new family name:

Anello is derived from latin “anello”, *the ring*, and relates to the circular nature of the DNA genome.

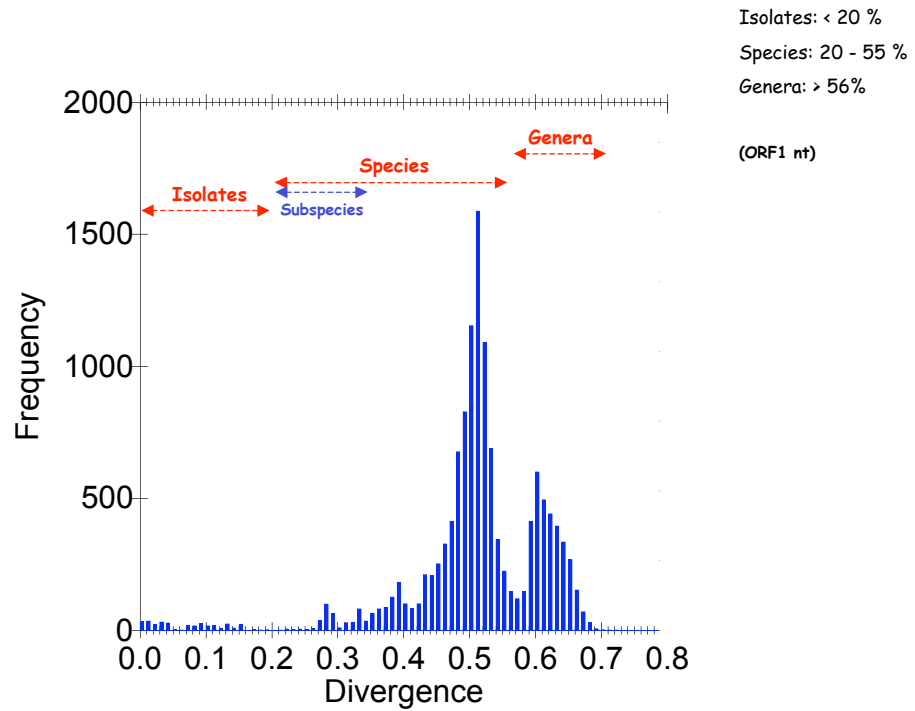
References:

- Biagini, P., Todd, D., Bendinelli, M., Hino, S., Mankertz, A., Mishiro, S., Niel, C., Okamoto, H., Raidal, S., Ritchie, B.W & Teo, G.C. (2005).** *Anellovirus*. In: *Virus Taxonomy*, VIIIth Report of the International Committee for the Taxonomy of Viruses (C.M. Fauquet, M.A. Mayo, J. Maniloff, U. Desselberger, and L.A. Ball, eds), 335-341. Elsevier/Academic Press, London.
- Biagini, P., Uch, R., Belhouchet, M., Attoui, H., Cantaloube, J.F., Brisbarre, N. & de Micco, P. (2007).** Circular genomes related to anelloviruses identified in human and animal samples using a combined rolling-circle amplification- sequence independent single primer amplification approach. *J Gen Virol* **88**, 2696-2701.
- Jelcic, I., Hotz-Wagenblatt, A., Hunziker, A., Zur Hausen, H. & de Villiers, E. M. (2004).** Isolation of multiple TT virus genotypes from spleen biopsy tissue from a Hodgkin's disease patient: genome reorganization and diversity in the hypervariable region. *J Virol* **78**, 7498-507.
- Jones, M. S., Kapoor, A., Lukashov, V. V., Simmonds, P., Hecht, F. & Delwart, E. (2005).** New DNA viruses identified in patients with acute viral infection syndrome. *J Virol* **79**, 8230-8236.
- Ninomiya, M., Nishizawa T., Takahashi, M., Lorenzo, F.R., Shimosegawa, T. & Okamoto, H. (2007).** Identification and genomic characterization of a novel human torque teno virus of 3.2 kb. *J Gen Virol* **88**, 1939-1944.
- Nishizawa, T., Okamoto, H., Konishi, K., Yoshizawa, H., Miyakawa, Y. & Mayumi, M. (1997).** A novel DNA virus (TTV) associated with elevated transaminase levels in posttransfusion hepatitis of unknown etiology. *Biochem Biophys Res Commun* **241**, 92-97.
- Okamoto, H., Nishizawa, T., Tawara, A., Peng, Y., Takahashi, M., Kishimoto, J., Tanaka, T., Miyakawa, Y. & Mayumi, M. (2000).** Species-specific TT viruses in humans and nonhuman primates and their phylogenetic relatedness. *Virology* **277**, 368-378.
- Okamoto, H., Takahashi, M., Nishizawa, T., Tawara, A., Fukai, K., Muramatsu, U., Naito, Y. & Yoshikawa, A. (2002).** Genomic characterization of TT viruses (TTVs) in pigs, cats and dogs and their relatedness with species-specific TTVs in primates and tupaia. *J Gen Virol* **83**, 1291-1297.
- Peng, Y. H., Nishizawa, T., Takahashi, M., Ishikawa, T., Yoshikawa, A. & Okamoto, H. (2002).** Analysis of the entire genomes of thirteen TT virus variants classifiable into the fourth and fifth genetic groups, isolated from viremic infants. *Arch Virol* **147**, 21-41.
- Takahashi, K., Iwasa, Y., Hijikata, M. & Mishiro, S. (2000).** Identification of a new human DNA virus (TTV-like mini virus, TLMV) intermediately related to TT virus and chicken anemia virus. *Arch Virol* **145**, 979-993.

Annexes:

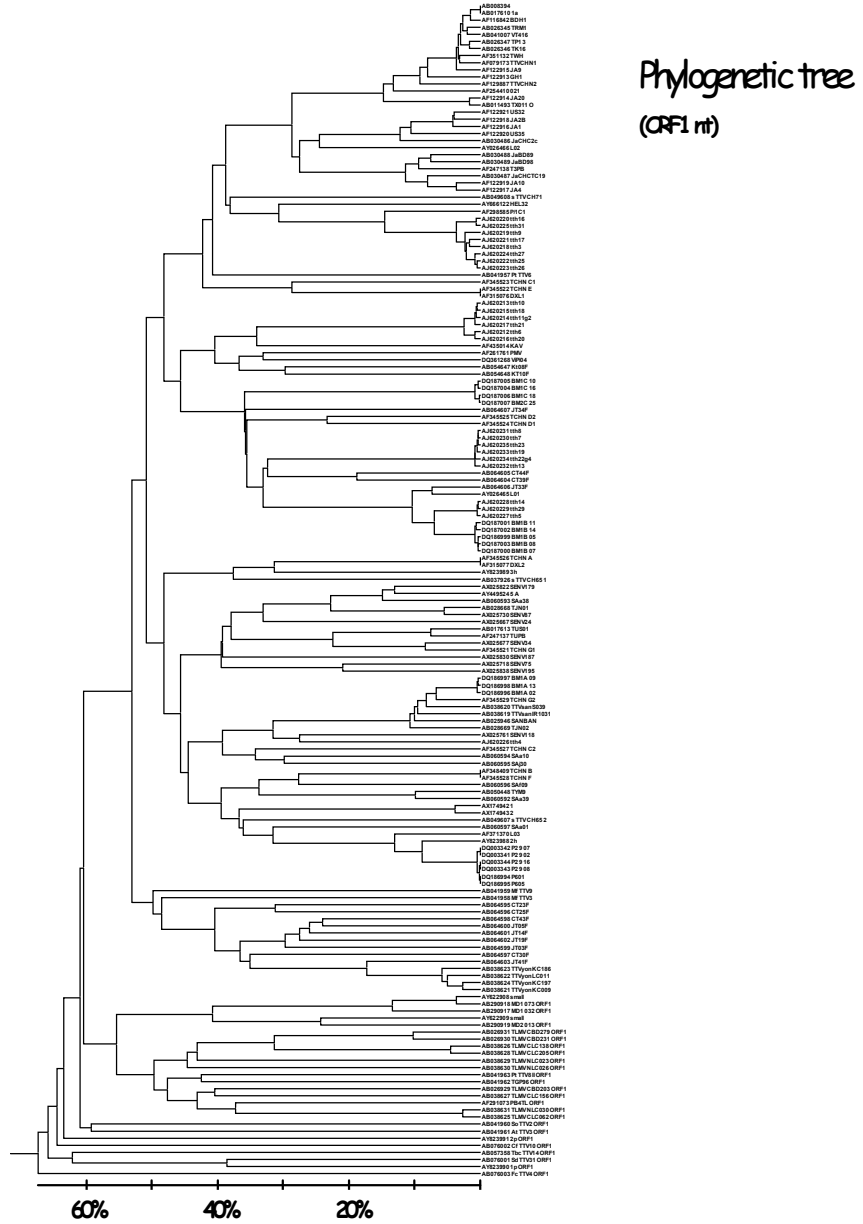
The classification has to reflect the extensive variability existing between members of the proposed family. The progressive discovery of highly divergent, complete *Anellovirus* genomes ranging from ~2 kb to ~4 kb in humans and other animals impairs a reliable phylogenetic and taxonomic analysis of full-length sequences. Based on these considerations, the analysis of the entire ORF1 (encoding the putative nucleocapsid protein) at the nucleotide level (ORF1-nt) is the most convenient approach. Analysis of the distribution of pairwise comparisons (Figure 1) and the corresponding phylogenetic tree (Figure 2) facilitates identification of the levels of genera, species and subspecies. Based on the currently available data, a taxonomic classification is proposed with the following cut-off values for sequence divergence: genera >56%, species >35%.

Figure 1. Analysis of the distribution of pairwise comparisons in ORF1-nt (166 sequences).



Distribution of pairwise comparisons

Figure 2. UPGMA phylogenetic tree built with ORF1-nt sequences.



ASSIGNMENT OF ACCESSION NUMBER TO TAXA

Alphatorquevirus

<i>Torque teno virus 1</i>	AB008394
<i>Torque teno virus 2</i>	AB049608
<i>Torque teno virus 3</i>	AY666122
<i>Torque teno virus 4</i>	AB041957
<i>Torque teno virus 5</i>	AF345523
<i>Torque teno virus 6</i>	AF435014
<i>Torque teno virus 7</i>	AF261761
<i>Torque teno virus 8</i>	AB054647
<i>Torque teno virus 9</i>	DQ187006
<i>Torque teno virus 10</i>	AB064607
<i>Torque teno virus 11</i>	AF345524
<i>Torque teno virus 12</i>	AB064605
<i>Torque teno virus 13</i>	AF345526
<i>Torque teno virus 14</i>	AB037926
<i>Torque teno virus 15</i>	AB028668
<i>Torque teno virus 16</i>	AB017613
<i>Torque teno virus 17</i>	AX025830
<i>Torque teno virus 18</i>	AX025718
<i>Torque teno virus 19</i>	AB025946
<i>Torque teno virus 20</i>	AB060594
<i>Torque teno virus 21</i>	AF348409
<i>Torque teno virus 22</i>	AX174942
<i>Torque teno virus 23</i>	AB049607
<i>Torque teno virus 24</i>	AB060597
<i>Torque teno virus 25</i>	AB041959
<i>Torque teno virus 26</i>	AB041958
<i>Torque teno virus 27</i>	AB064595
<i>Torque teno virus 28</i>	AB064598
<i>Torque teno virus 29</i>	AB038621

Betatorquevirus

<i>Torque teno mini virus 1</i>	AB026931
<i>Torque teno mini virus 2</i>	AB038629
<i>Torque teno mini virus 3</i>	AB038630
<i>Torque teno mini virus 4</i>	AB041963
<i>Torque teno mini virus 5</i>	AB041962
<i>Torque teno mini virus 6</i>	AB026929
<i>Torque teno mini virus 7</i>	AB038627
<i>Torque teno mini virus 8</i>	AF291073
<i>Torque teno mini virus 9</i>	AB038631

Gammatorquevirus

<i>Torque teno midi virus 1</i>	AB290918
<i>Torque teno midi virus 2</i>	AB290919

Deltatorquevirus

<i>Torque teno tupaia virus</i>	AB057358
---------------------------------	----------

Epsilontorquevirus

<i>Torque teno tamarin virus</i>	AB041960
----------------------------------	----------

Zetatorquevirus

<i>Torque teno douroucouli virus</i>	AB041961
--------------------------------------	----------

Etatorquevirus

<i>Torque teno felis virus</i>	AB076003
--------------------------------	----------

Thetatorquevirus

<i>Torque teno canis virus</i>	AB076002
--------------------------------	----------

Iotatorquevirus

<i>Torque teno sus virus 1</i>	AB076001
<i>Torque teno sus virus 2</i>	AY823990

NB: The nucleotide sequence AY823991 also identified in swine presents an extreme sequence divergence compatible with the creation of a tenth genus. However, it is proposed to keep it as an unassigned virus in the genus *Iotatorquevirus* until further data have been collected in swine species.

MODULE 4: **NEW GENUS**

Code	<i>2007.075dV</i>	(assigned by ICTV officers)
To create a new genus assigned as follows:		
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Code	<i>2007.075eV</i>	(assigned by ICTV officers)
To name the new genus: <i>Alphatorquevirus</i>		

Code	<i>2007.075fV</i>	(assigned by ICTV officers)
To assign the following as species in the new genus:		
<i>Torque teno virus 1</i>		
<i>Torque teno virus 2</i>		
<i>Torque teno virus 3</i>		
<i>Torque teno virus 4</i>		
<i>Torque teno virus 5</i>		
<i>Torque teno virus 6</i>		
<i>Torque teno virus 7</i>		
<i>Torque teno virus 8</i>		
<i>Torque teno virus 9</i>		
<i>Torque teno virus 10</i>		
<i>Torque teno virus 11</i>		
<i>Torque teno virus 12</i>		
<i>Torque teno virus 13</i>		
<i>Torque teno virus 14</i>		
<i>Torque teno virus 15</i>		
<i>Torque teno virus 16</i>		
<i>Torque teno virus 17</i>		
<i>Torque teno virus 18</i>		
<i>Torque teno virus 19</i>		
<i>Torque teno virus 20</i>		
<i>Torque teno virus 21</i>		
<i>Torque teno virus 22</i>		
<i>Torque teno virus 23</i>		
<i>Torque teno virus 24</i>		
<i>Torque teno virus 25</i>		
<i>Torque teno virus 26</i>		
<i>Torque teno virus 27</i>		
<i>Torque teno virus 28</i>		
<i>Torque teno virus 29</i>		
[All new]		

Code	2007.075gV	(assigned by ICTV officers)
To designate the following as the type species in the new genus:		
<i>Torque teno virus 1</i>		

Argument to justify the creation of a new genus:

See Module 2.

Origin of the new genus name:

Torque is derived from latin “torques”, *the necklace*, and relates to the circular, single-stranded nature of the DNA genome. *Teno* is derived from latin “tenuis”, *thin*, and relates to the small size of the viral genome.

Argument to justify the choice of type species:

First numbered in series.

Species demarcation criteria in the genus:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 4: **NEW GENUS**

Code	<i>2007.075hV</i>	(assigned by ICTV officers)
To create a new genus assigned as follows:		
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Code	<i>2007.075iV</i>	(assigned by ICTV officers)
To name the new genus: <i>Betatorquevirus</i>		

Code	<i>2007.075jV</i>	(assigned by ICTV officers)
To assign the following as species in the new genus:		
<i>Torque teno mini virus 1</i>		
<i>Torque teno mini virus 2</i>		
<i>Torque teno mini virus 3</i>		
<i>Torque teno mini virus 4</i>		
<i>Torque teno mini virus 5</i>		
<i>Torque teno mini virus 6</i>		
<i>Torque teno mini virus 7</i>		
<i>Torque teno mini virus 8</i>		
<i>Torque teno mini virus 9</i>		
[All new]		

Code	<i>2007.075kV</i>	(assigned by ICTV officers)
To designate the following as the type species in the new genus:		
<i>Torque teno mini virus 1</i>		

Argument to justify the creation of a new genus:

See Module 2.

Origin of the new genus name:

Torque is derived from latin “torques”, *the necklace*, and relates to the circular, single-stranded nature of the DNA genome. *Teno* is derived from latin “tenuis”, *thin*, and relates to the small size of the viral genome. *Mini* is derived from latin “minimus”, *small*, and relates to the smaller size of the genomes related phylogenetically, when compared to those of *Torque teno virus* members.

Argument to justify the choice of type species:

First numbered in series.

Species demarcation criteria in the genus:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 4: **NEW GENUS**

Code	<i>2007.075lV</i>	(assigned by ICTV officers)
To create a new genus assigned as follows:		
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Code	<i>2007.075mV</i>	(assigned by ICTV officers)
To name the new genus: <i>Gammatorquevirus</i>		

Code	<i>2007.075nV</i>	(assigned by ICTV officers)
To assign the following as species in the new genus:		
<i>Torque teno midi virus 1</i>		
<i>Torque teno midi virus 2</i>		
[Both new]		

Code	<i>2007.075oV</i>	(assigned by ICTV officers)
To designate the following as the type species in the new genus:		
<i>Torque teno midi virus 1</i>		

Argument to justify the creation of a new genus:

See Module 2.

Origin of the new genus name:

Torque is derived from latin “torques”, *the necklace*, and relates to the circular, single-stranded nature of the DNA genome. *Teno* is derived from latin “tenuis”, *thin*, and relates to the small size of the viral genome. *Midi* is derived from latin “medius”, *intermediate*, and relates to the intermediate size of the genomes related phylogenetically, when compared to those of *Torque teno virus* and *Torque teno mini virus* members.

Argument to justify the choice of type species:

First numbered in series.

Species demarcation criteria in the genus:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 4: **NEW GENUS**

Code	<i>2007.075pV</i>	(assigned by ICTV officers)
To create a new genus assigned as follows:		
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Code	<i>2007.075qV</i>	(assigned by ICTV officers)
To name the new genus: <i>Deltatorquevirus</i>		

Code	<i>2007.075rV</i>	(assigned by ICTV officers)
To assign the following as species in the new genus:		
<i>Torque teno tupaia virus</i>		
[New]		

Code	<i>2007.075sV</i>	(assigned by ICTV officers)
To designate the following as the type species in the new genus:		
<i>Torque teno tupaia virus</i>		

Argument to justify the creation of a new genus:

See Module 2.

Origin of the new genus name:

Torque is derived from latin “torques”, *the necklace*, and relates to the circular, single-stranded nature of the DNA genome. *Teno* is derived from latin “tenuis”, *thin*, and relates to the small size of the viral genome. *Tupaia* relates to the animal species in which the virus was first identified.

Argument to justify the choice of type species:

First species in the genus.

Species demarcation criteria in the genus:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 4: **NEW GENUS**

(if more than one genus is to be created, please complete additional copies of this section)

Code	<i>2007.075tV</i>	(assigned by ICTV officers)
To create a new genus assigned as follows:		
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Code	<i>2007.075uV</i>	(assigned by ICTV officers)
To name the new genus: <i>Epsilontorquevirus</i>		

Code	<i>2007.075vV</i>	(assigned by ICTV officers)
To assign the following as species in the new genus:		
<i>Torque teno tamarin virus</i>		
[New]		

Code	<i>2007.075wV</i>	(assigned by ICTV officers)
To designate the following as the type species in the new genus:		
<i>Torque teno tamarin virus</i>		

Argument to justify the creation of a new genus:

See Module 2.

Origin of the new genus name:

Torque is derived from latin “torques”, *the necklace*, and relates to the circular, single-stranded nature of the DNA genome. *Teno* is derived from latin “tenuis”, *thin*, and relates to the small size of the viral genome. *Tamarin* relates to the animal species in which the virus was first identified.

Argument to justify the choice of type species:

First species in the genus.

Species demarcation criteria in the genus:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 4: **NEW GENUS**

Code	2007.075xV	(assigned by ICTV officers)
To create a new genus assigned as follows:		
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Code	2007.075yV	(assigned by ICTV officers)
To name the new genus: <i>Zetatorquevirus</i>		

Code	2007.075zV	(assigned by ICTV officers)
To assign the following as species in the new genus:		
<i>Torque teno douroucouli virus</i>		
[New]		

Code	2007.075aaV	(assigned by ICTV officers)
To designate the following as the type species in the new genus:		
<i>Torque teno douroucouli virus</i>		

Argument to justify the creation of a new genus:

See Module 2.

Origin of the new genus name:

Torque is derived from latin “torques”, *the necklace*, and relates to the circular, single-stranded nature of the DNA genome. *Teno* is derived from latin “tenuis”, *thin*, and relates to the small size of the viral genome. *Douroucouli* relates to the animal species in which the virus was first identified.

Argument to justify the choice of type species:

First species in the genus.

Species demarcation criteria in the genus:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 4: **NEW GENUS**

Code	<i>2007.075bbV</i>	(assigned by ICTV officers)
To create a new genus assigned as follows:		
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Code	<i>2007.075ccV</i>	(assigned by ICTV officers)
To name the new genus: <i>Etatorquevirus</i>		

Code	<i>2007.075ddV</i>	(assigned by ICTV officers)
To assign the following as species in the new genus:		
<i>Torque teno felis virus</i>		
[New]		

Code	<i>2007.075eeV</i>	(assigned by ICTV officers)
To designate the following as the type species in the new genus:		
<i>Torque teno felis virus</i>		

Argument to justify the creation of a new genus:

See Module 2.

Origin of the new genus name:

Torque is derived from latin “torques”, *the necklace*, and relates to the circular, single-stranded nature of the DNA genome. *Teno* is derived from latin “tenuis”, *thin*, and relates to the small size of the viral genome. *Felis* relates to the animal species in which the virus was first identified.

Argument to justify the choice of type species:

First species in the genus.

Species demarcation criteria in the genus:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 4: **NEW GENUS**

Code	<i>2007.075ffV</i>	(assigned by ICTV officers)
To create a new genus assigned as follows:		
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Code	<i>2007.075ggV</i>	(assigned by ICTV officers)
To name the new genus: <i>Thetatorquevirus</i>		

Code	<i>2007.075hhV</i>	(assigned by ICTV officers)
To assign the following as species in the new genus:		
<i>Torque teno canis virus</i>		
[New]		

Code	<i>2007.075iiV</i>	(assigned by ICTV officers)
To designate the following as the type species in the new genus:		
<i>Torque teno canis virus</i>		

Argument to justify the creation of a new genus:

See Module 2.

Origin of the new genus name:

Torque is derived from latin “torques”, *the necklace*, and relates to the circular, single-stranded nature of the DNA genome. *Teno* is derived from latin “tenuis”, *thin*, and relates to the small size of the viral genome. *Canis* relates to the animal species in which the virus was first identified.

Argument to justify the choice of type species:

First species in the genus.

Species demarcation criteria in the genus:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 4: **NEW GENUS**

(if more than one genus is to be created, please complete additional copies of this section)

Code	<i>2007.075jjV</i>	(assigned by ICTV officers)
To create a new genus assigned as follows:		
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Code	<i>2007.075kkV</i>	(assigned by ICTV officers)
To name the new genus: <i>Iotatorquevirus</i>		

Code	<i>2007.075lUV</i>	(assigned by ICTV officers)
To assign the following as species in the new genus:		
<i>Torque teno sus virus 1</i>		
<i>Torque teno sus virus 2</i>		
[Both new]		

Code	<i>2007.075mmV</i>	(assigned by ICTV officers)
To designate the following as the type species in the new genus:		
<i>Torque teno sus virus 1</i>		

Argument to justify the creation of a new genus:

See Module 2.

Origin of the new genus name:

Torque is derived from latin “torques”, *the necklace*, and relates to the circular, single-stranded nature of the DNA genome. *Teno* is derived from latin “tenuis”, *thin*, and relates to the small size of the viral genome. *Sus* relates to the animal species in which the viruses were first identified.

Argument to justify the choice of type species:

First numbered in series.

Species demarcation criteria in the genus:

See Module 2.

References:

References:

See Module 2.

Annexes:

See Module 2.

MODULE 5: **NEW SPECIES**

Code	2007.075nnV	(assigned by ICTV officers)
To create 29 new species assigned as follows:		
Genus:	<i>Alphatorquevirus</i>	
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Name(s) of proposed new species:

- Torque teno virus 1*
- Torque teno virus 2*
- Torque teno virus 3*
- Torque teno virus 4*
- Torque teno virus 5*
- Torque teno virus 6*
- Torque teno virus 7*
- Torque teno virus 8*
- Torque teno virus 9*
- Torque teno virus 10*
- Torque teno virus 11*
- Torque teno virus 12*
- Torque teno virus 13*
- Torque teno virus 14*
- Torque teno virus 15*
- Torque teno virus 16*
- Torque teno virus 17*
- Torque teno virus 18*
- Torque teno virus 19*
- Torque teno virus 20*
- Torque teno virus 21*
- Torque teno virus 22*
- Torque teno virus 23*
- Torque teno virus 24*
- Torque teno virus 25*
- Torque teno virus 26*
- Torque teno virus 27*
- Torque teno virus 28*
- Torque teno virus 29*

Argument to justify the creation of the new species:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 5: **NEW SPECIES**

Code	2007.07500V	(assigned by ICTV officers)
To create 9 new species assigned as follows:		
Genus:	<i>Betatorquevirus</i>	
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Name(s) of proposed new species:

<i>Torque teno mini virus 1</i> <i>Torque teno mini virus 2</i> <i>Torque teno mini virus 3</i> <i>Torque teno mini virus 4</i> <i>Torque teno mini virus 5</i> <i>Torque teno mini virus 6</i> <i>Torque teno mini virus 7</i> <i>Torque teno mini virus 8</i> <i>Torque teno mini virus 9</i>

Argument to justify the creation of the new species:

|
See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 5: **NEW SPECIES**

Code	<i>2007.075ppV</i>	(assigned by ICTV officers)
To create 1 new species assigned as follows:		
Genus:	<i>Gammatorquevirus</i>	
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Name(s) of proposed new species:

<i>Torque teno midi virus 1</i> <i>Torque teno midi virus 2</i>
--

Argument to justify the creation of the new species:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 5: **NEW SPECIES**

Code	<i>2007.075qqV</i>	(assigned by ICTV officers)
To create 1 new species assigned as follows:		
Genus:	<i>Deltatorquevirus</i>	
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Name(s) of proposed new species:

Torque teno tupaia virus

Argument to justify the creation of the new species:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 5: **NEW SPECIES**

Code	<i>2007.075rrV</i>	(assigned by ICTV officers)
To create 1 new species assigned as follows:		
Genus:	<i>Epsilontorquevirus</i>	
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Name(s) of proposed new species:

Torque teno tamarin virus

Argument to justify the creation of the new species:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 5: **NEW SPECIES**

Code	2007.075ssV	(assigned by ICTV officers)
To create 1 new species assigned as follows:		
Genus:	<i>Zetatorquevirus</i>	
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Name(s) of proposed new species:

Torque teno douroucouli virus

Argument to justify the creation of the new species:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 5: **NEW SPECIES**

Code	2007.075ttV	(assigned by ICTV officers)
To create 1 new species assigned as follows:		
Genus:	<i>Etatorquevirus</i>	
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Name(s) of proposed new species:

Torque teno felis virus

Argument to justify the creation of the new species:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 5: **NEW SPECIES**

Code	<i>2007.07uu5V</i>	(assigned by ICTV officers)
To create 1 new species assigned as follows:		
Genus:	<i>Thetatorquevirus</i>	
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Name(s) of proposed new species:

Torque teno canis virus

Argument to justify the creation of the new species:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 5: **NEW SPECIES**

Code	2007.075vvV	(assigned by ICTV officers)
To create 2 new species assigned as follows:		
Genus:	<i>Iotatorquevirus</i>	
Subfamily:	<i>Unassigned</i>	
Family:	<i>Anelloviridae</i>	
Order:	<i>Unassigned</i>	

Name(s) of proposed new species:

<i>Torque teno sus virus 1</i> <i>Torque teno sus virus 2</i>
--

Argument to justify the creation of the new species:

See Module 2.

References:

See Module 2.

Annexes:

See Module 2.

MODULE 6: **REMOVE and MOVE**

SECTION (a)

Code	<i>2007.075wwV</i>	(assigned by ICTV officers)
To remove (abolish) the following taxon(s):		
<i>Anellovirus</i>		

Old and new composition of the higher taxon that will be depleted by the removal:

No higher taxon.

Argument to justify the removal:

Creation of new family, new genera and new species.

Code	<i>2007.075xxV</i>	(assigned by ICTV officers)
To remove (abolish) the following taxon(s):		
<i>Torque teno virus</i>		

Old and new composition of the higher taxon that will be depleted by the removal:

Genus <i>Anellovirus</i> and one species therein removed. Genus promoted to new family. Single species replaced by 47 new species.
--

Argument to justify the removal:

Creation of new family, new genera and new species.
