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20	Rhabdoviridae; Sunviridae; virus classification; virus nomenclature; virus taxonomy

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### 1 ABSTRACT

In 2016, the order *Mononegavirales* was emended through the addition of two new families
(*Mymonaviridae* and *Sunviridae*), the elevation of the paramyxoviral subfamily *Pneumovirinae*to family status (*Pneumoviridae*), the addition of five free-floating genera (*Anphevirus, Arlivirus, Chengtivirus, Crustavirus*, and *Wastrivirus*), and several other changes at the genus and species
levels. This article presents the updated taxonomy of the order *Mononegavirales* as now accepted
by the International Committee on Taxonomy of Viruses (ICTV).

### 8 INTRODUCTION

The viral order *Mononegavirales* was established in 1991 to accommodate related viruses with 9 nonsegmented, linear, single-stranded negative-sense RNA genomes. These viruses were initially 10 11 assigned to three mononegaviral families: *Filoviridae*, *Paramyxoviridae*, and *Rhabdoviridae* [20, 21]. In subsequent years, these families continued to grow through the inclusion of numerous 12 novel species and genera, and the order was therefore emended in 1995 [4], 1997 [23], 2000 13 14 [24], 2005 [25], and 2011 [8]. The families *Bornaviridae* and *Nyamiviridae* joined the other three mononegaviral families in 1996 [22] and 2014 [1, 11], respectively. In 2015, the Study Groups 15 of the International Committee on Taxonomy of Viruses (ICTV) responsible for the taxonomy of 16 the order and its five families embarked on a joint effort to assign unclassified mononegaviruses 17 to existing or novel taxa and to streamline order nomenclature. Here we present a brief overview 18 of the first round of these efforts, which by now is accepted by the ICTV Executive Committee 19 and, thereby, is official taxonomy. 20

# 21 TAXONOMIC CHANGES AT THE ORDER LEVEL

1	In recent years, several mononegaviruses have been described that are only distantly related to
2	the members of the families Bornaviridae, Filoviridae, Nyamiviridae, Paramyxoviridae, and
3	Rhabdoviridae. These viruses include Sclerotinia sclerotiorum negative-stranded RNA virus 1
4	(SsNSRV-1) found in an ascomycete in China [16]; Sunshine Coast virus (SunCV; previously
5	called Sunshine virus) isolated from Australian carpet pythons [10]; and Líshí spider virus 2
6	(LsSV-2), Sānxiá water strider virus 4 (SxWSV-4), Tăchéng tick virus 6 (TcTV-6), Wēnzhōu
7	crab virus 1 (WzCV-1), and Xīnchéng mosquito virus (XcMV) detected in Chinese arthropods
8	[15]. To accommodate these viruses in the order and to appropriately reflect their phylogenetic
9	relationships to other mononegaviral taxa, two new families and four free-floating genera were
10	established: Mymonaviridae (accommodating SsNSRV-1), Sunviridae (SunCV), Anphevirus
11	(XcMV), Arlivirus (LsSV-2), Chengtivirus (TcTV-6), Crustavirus (WzCV-1), and Wastrivirus
12	(SxWSV-4). In addition, the paramyxoviral subfamily Pneumovirinae was elevated to family
13	status (Pneumoviridae) because the members of this taxon are as closely related to filoviruses as
14	to the members of the paramyxoviral subfamily Paramxyovirinae (now dissolved) (Table 1).

# 15 TAXONOMIC CHANGES AT THE FAMILY LEVEL

The monogeneric family *Bornaviridae* was reorganized in 2015 by establishing five distinct
species in the genus *Bornavirus* [2, 12] following a non-Latinized binomial species name format
[29]. These efforts were continued in 2016 by expanding the genus by an additional two species
(*Elapid 1 bornavirus* for Loveridge's garter snake virus 1 [27] and *Psittaciform 2 bornavirus* for
parrot bornavirus 5 [9, 18]) (Table 1).

The monogeneric family *Nyamiviridae* was expanded to include a second genus (*Socyvirus*) for
 the until-then free-floating nyamivirus species *Soybean cyst nematode virus*. This species name
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was changed to *Soybean cyst nematode socyvirus* to adhere to the non-Latinized binomial
 species name format [29] (Table 1).

3	The family Paramyxoviridae was emended by expanding the genus Avulavirus by three species
4	(Avian paramyxovirus 10-12 for avian paramyxoviruses 10-12, respectively [5, 19, 28]), the
5	genus Henipavirus by three species (Cedar henipavirus for Cedar virus [17], Ghanaian bat
6	henipavirus for Kumasi virus [GH-M74a] [7], and Mojiang henipavirus for Mojiang virus [31]),
7	the genus <i>Morbillivirus</i> by one species ( <i>Feline morbillivirus</i> for feline morbillivirus [30]) and the
8	genus Respirovirus by one species (Porcine parainfluenza virus 1 for porcine parainfluenza virus
9	1 [14]). The species Simian Virus 10 was dissolved on the evidence that simian virus 10 is an
10	isolate of human parainfluenzavirus 3 rather than a distinct virus [13]. The genus <i>Pneumovirus</i> ,
11	now included in the new family Pneumoviridae, was renamed Orthopneumovirus to avoid
12	confusion between family and genus members (Table 1).
13	The family <i>Rhabdoviridae</i> was expanded by two genera: <i>Dichorhavirus</i> (new; [6]) and
14	Varicosavirus (previously free-floating outside of the order) to accommodate bisegmented plant
15	viruses (coffee ringspot virus and orchid fleck virus; lettuce big-vein associated virus). The
16	species Alfalfa dwarf cytorhabdovirus (for alfalfa dwarf virus [3]) was added to the genus
17	Cytorhabdovirus. Finally, the non-Latinized binomial species name format [29] was applied
18	$(1, \dots, 1, 1, \dots, 1) \rightarrow (T_1, 1)$
	throughout the family (Table 1).

in Table 1.

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# **Table 1. Taxonomy of the order** *Mononegavirales* **as of 2016.** Listed are all mononegaviruses that have been classified into species.

# 2 Asterisks denote type species.

Genus	Species	Virus (Abbreviation)
	Family Bornaviridae	I
Bornavirus	Elapid 1 bornavirus	Loveridge's garter snake virus 1 (LGSV-1)
	Mammalian 1 bornavirus*	Borna disease virus 1 (BoDV-1)
		Borna disease virus 2 (BoDV-2)
	Passeriform 1 bornavirus	canary bornavirus 1 (CnBV-1)
		canary bornavirus 2 (CnBV-2)
		canary bornavirus 3 (CnBV-3)
	Passeriform 2 bornavirus	estrildid finch bornavirus 1 (EsBV-1)
	Psittaciform 1 bornavirus	parrot bornavirus 1 (PaBV-1)
		parrot bornavirus 2 (PaBV-2)
		parrot bornavirus 3 (PaBV-3)
		parrot bornavirus 4 (PaBV-4)
		parrot bornavirus 7 (PaBV-7)

Psittaciform 2 bornavirus	parrot bornavirus 5 (PaBV-5)
Waterbird 1 bornavirus	aquatic bird bornavirus 1 (ABBV-1)
	aquatic bird bornavirus 2 (ABBV-2)
Family <i>Filoviridae</i>	
Lloviu cuevavirus*	Lloviu virus (LLOV)
Bundibugyo ebolavirus	Bundibugyo virus (BDBV)
Reston ebolavirus	Reston virus (RESTV)
Sudan ebolavirus	Sudan virus (SUDV)
Taï Forest ebolavirus	Taï Forest virus (TAFV)
Zaire ebolavirus*	Ebola virus (EBOV)
Marburg marburgvirus*	Marburg virus (MARV)
	Ravn virus (RAVV)
Family Mymonaviridae	
Sclerotinia sclerotimonavirus*	Sclerotinia sclerotiorum negative-stranded
	RNA virus 1 (SsNSRV-1)
	Waterbird 1 bornavirus         Waterbird 1 bornavirus         Family Filoviridae         Lloviu cuevavirus*         Bundibugyo ebolavirus         Bundibugyo ebolavirus         Reston ebolavirus         Sudan ebolavirus         Taï Forest ebolavirus         Zaire ebolavirus*         Marburg marburgvirus*         Family Mymonaviridae

Family Nyamiviridae		
Nyavirus	Midway nyavirus	Midway virus (MIDWV)
	Nyamanini nyavirus*	Nyamanini virus (NYMV)
	Sierra Nevada nyavirus	Sierra Nevada virus (SNVV)
Socyvirus	Soybean cyst nematode socyvirus*	soybean cyst nematode virus 1 (SbCNV-1)
	Family Paramyxoviridae	
Aquaparamyxovirus	Atlantic salmon paramyxovirus*	Atlantic salmon paramyxovirus (AsaPV)
Avulavirus	Avian paramyxovirus 2	avian paramyxovirus 2 (APMV-2)
	Avian paramyxovirus 3	avian paramyxovirus 3 (APMV-3)
	Avian paramyxovirus 4	avian paramyxovirus 4 (APMV-4)
	Avian paramyxovirus 5	avian paramyxovirus 5 (APMV-5)
	Avian paramyxovirus 6	avian paramyxovirus 6 (APMV-6)
	Avian paramyxovirus 7	avian paramyxovirus 7 (APMV-7)
	Avian paramyxovirus 8	avian paramyxovirus 8 (APMV-8)
	Avian paramyxovirus 9	avian paramyxovirus 9 (APMV-9)

	Avian paramyxovirus 10	avian paramyxovirus 10 (APMV-10)
	Avian paramyxovirus 11	avian paramyxovirus 11 (APMV-11)
	Avian paramyxovirus 12	avian paramyxovirus 12 (APMV-12)
	Newcastle disease virus*	avian paramyxovirus 1 (APMV-1) <sup>1</sup>
Ferlavirus	Fer-de-Lance paramyxovirus*	Fer-de-Lance virus (FDLV) <sup>2</sup>
Henipavirus	Cedar henipavirus	Cedar virus (CedV)
	Ghanaian bat henipavirus	Kumasi virus (KV) <sup>3</sup>
	Hendra virus*	Hendra virus (HeV)
	Mojiang henipavirus	Mòjiāng virus (MojV)
	Nipah virus	Nipah virus (NiV)
Morbillivirus	Canine distemper virus	canine distemper virus (CDV)
	Cetacean morbillivirus	cetacean morbillivirus (CeMV)
	Feline morbillivirus	feline morbillivirus (FeMV) <sup>4</sup>
	Measles virus*	measles virus (MeV)
	Peste-des-petits-ruminants virus	peste-des-petits-ruminants virus (PPRV)

	Phocine distemper virus	phocine distemper virus (PDV)
	Rinderpest virus	rinderpest virus (RPV)
Respirovirus	Bovine parainfluenza virus 3	bovine parainfluenza virus 3 (BPIV-3)
	Human parainfluenza virus 1	human parainfluenza virus 1 (HPIV-1)
	Human parainfluenza virus 3	human parainfluenza virus 3 (HPIV-3) <sup>5</sup>
	Porcine parainfluenza virus 1	porcine parainfluenza virus 1 (PPIV-1)
	Sendai virus*	Sendai virus (SeV) <sup>6</sup>
Rubulavirus	Human parainfluenza virus 2	human parainfluenza virus 2 (HPIV-2)
	Human parainfluenza virus 4	human parainfluenza virus 4a (HPIV-4a)
		human parainfluenza virus 4b (HPIV-4b)
	Mapuera virus	Mapuera virus (MapV)
	Mumps virus*	mumps virus (MuV)
		bat mumps virus (BMV) <sup>7</sup>
	Parainfluenza virus 5	parainfluenza virus 5 (PIV-5) <sup>8</sup>
	Porcine rubulavirus	La Piedad Michoacán Mexico virus

	(LPMV) <sup>9</sup>
Simian virus 41	simian virus 41 (SV-41)
Family Pneumoviridae	
Avian metapneumovirus*	avian metapneumovirus (AMPV) <sup>10</sup>
Human metapneumovirus	human metapneumovirus (HMPV)
Bovine respiratory syncytial virus	bovine respiratory syncytial virus (BRSV)
Human respiratory syncytial virus*	human respiratory syncytial virus A2
	(HRSV-A2)
	human respiratory syncytial virus B1
	(HRSV-B1)
	human respiratory syncytial virus S2
	(HRSV-S2)
Murine pneumonia virus	murine pneumonia virus (MPV)
Family <i>Rhabdoviridae</i>	
Alfalfa dwarf cytorhabdovirus	alfalfa dwarf virus (ADV)
Barley yellow striate mosaic	barley yellow striate mosaic virus
	Family Pneumoviridae         Family Pneumoviridae         Avian metapneumovirus*         Human metapneumovirus         Bovine respiratory syncytial virus*         Human respiratory syncytial virus*         Human respiratory syncytial virus*         Murine pneumonia virus         Family Rhabdoviridae         Alfalfa dwarf cytorhabdovirus

	cytorhabdovirus	(BYSMV)
	Broccoli necrotic yellows cytorhabdovirus	broccoli necrotic yellows virus (BNYV)
	Festuca leaf streak cytorhabdovirus	festuca leaf streak virus (FLSV)
	Lettuce necrotic yellows cytorhabdovirus*	lettuce necrotic yellows virus (LNYV)
	Lettuce yellow mottle cytorhabdovirus	lettuce yellow mottle virus (LYMoV)
	Northern cereal mosaic cytorhabdovirus	northern cereal mosaic virus (NCMV)
	Sonchus cytorhabdovirus 1	sonchus virus (SonV)
	Strawberry crinkle cytorhabdovirus	strawberry crinkle virus (SCV)
	Wheat American striate mosaic	wheat American striate mosaic virus
	cytorhabdovirus	(WASMV)
Dichorhavirus	Coffee ringspot dichorhavirus	coffee ringspot virus (CoRSV)
	Orchid fleck dichorhavirus*	orchid fleck virus (OFV) <sup>11</sup>
Ephemerovirus	Adelaide River ephemerovirus	Adelaide River virus (ARV)
	Berrimah ephemerovirus	Berrimah virus (BRMV)
	Bovine fever ephemerovirus*	bovine ephemeral fever virus (BEFV) <sup>12</sup>

	Kotonkan ephemerovirus	kotonkan virus (KOTV)
	Obodhiang ephemerovirus	Obodhiang virus (OBOV)
Lyssavirus	Aravan lyssavirus	Aravan virus (ARAV)
	Australian bat lyssavirus	Australian bat lyssavirus (ABLV)
	Bokeloh bat lyssavirus	Bokeloh bat lyssavirus (BBLV)
	Duvenhage lyssavirus	Duvenhage virus (DUVV)
	European bat 1 lyssavirus	European bat lyssavirus 1 (EBLV-1)
	European bat 2 lyssavirus	European bat lyssavirus 2 (EBLV-2)
	Ikoma lyssavirus	Ikoma lyssavirus (IKOV)
	Irkut lyssavirus	Irkut virus (IRKV)
	Khujand lyssavirus	Khujand virus (KHUV)
	Lagos bat lyssavirus	Lagos bat virus (LBV)
	Mokola lyssavirus	Mokola virus (MOKV)
	Rabies lyssavirus*	rabies virus (RABV)
	Shimoni bat lyssavirus	Shimoni bat virus (SHIBV)

	West Caucasian bat lyssavirus	West Caucasian bat virus (WCBV)
Novirhabdovirus	Hirame novirhabdovirus	Hirame rhabdovirus (HIRV)
	Oncorhynchus 1 novirhabdovirus*	infectious hematopoietic necrosis virus
		(IHNV)
	Oncorhynchus 2 novirhabdovirus	viral hemorrhagic septicemia virus
		(VHSV) <sup>13</sup>
	Snakehead novirhabdovirus	snakehead rhabdovirus (SHRV)
Nucleorhabdovirus	Datura yellow vein nucleorhabdovirus	datura yellow vein virus (DYVV)
	Eggplant mottled dwarf	eggplant mottled dwarf virus (EMDV)
	nucleorhabdovirus	
	Maize fine streak nucleorhabdovirus	maize fine streak virus (MSFV)
	Maize Iranian mosaic nucleorhabdovirus	maize Iranian mosaic virus (MIMV)
	Maize mosaic nucleorhabdovirus	maize mosaic virus (MMV)
	Potato yellow dwarf nucleorhabdovirus*	potato yellow dwarf virus (PYDV)
	Rice yellow stunt nucleorhabdovirus	rice yellow stunt virus (RYSV)
		rice transitory yellowing virus (RTYV)

Sonchus yellow net nucleorhabdovirus	sonchus yellow net virus (SYNV)
Sowthistle yellow vein nucleorhabdovirus	sowthistle yellow vein virus (SYVV)
Taro vein chlorosis nucleorhabdovirus	taro vein chlorosis virus (TaVCV)
Anguillid perhabdovirus	eel virus European X (EVEX)
Perch perhabdovirus*	perch rhabdovirus (PRV)
Sea trout perhabdovirus	lake trout rhabdovirus (LTRV)
Drosophila affinis sigmavirus	Drosophila affinis sigmavirus (DAffSV)
Drosophila ananassae sigmavirus	Drosophila ananassae sigmavirus
	(DAnaSV)
Drosophila immigrans sigmavirus	Drosophila immigrans sigmavirus
	(DImmSV)
Drosophila melanogaster sigmavirus*	Drosophila melanogaster sigmavirus
	(DMelSV)
Drosophila obscura sigmavirus	Drosophila obscura sigmavirus (DObsSV)
Drosophila tristis sigmavirus	Drosophila tristis sigmavirus (DTriSV
Muscina stabulans sigmavirus	Muscina stabulans sigmavirus (MStaSV)
	Sowthistle yellow vein nucleorhabdovirus         Taro vein chlorosis nucleorhabdovirus         Anguillid perhabdovirus         Perch perhabdovirus*         Sea trout perhabdovirus         Drosophila affinis sigmavirus         Drosophila immigrans sigmavirus         Drosophila melanogaster sigmavirus*         Drosophila obscura sigmavirus         Drosophila tristis sigmavirus

Sprivivirus	Carp sprivivirus*	spring viremia of carp virus (SVCV)
	Pike fry sprivivirus	grass carp rhabdovirus (GrCRV)
		pike fry rhabdovirus (PFRV)
		Tench rhabdovirus (TenRV)
Tibrovirus	Coastal Plains tibrovirus	Coastal Plains virus (CPV)
	Tibrogargan tibrovirus*	Bivens Arm virus (BAV)
		Tibrogargan virus (TIBV)
Tupavirus	Durham tupavirus*	Durham virus (DURV)
	Tupaia tupavirus	tupaia virus (TUPV)
Varicosavirus	Lettuce big-vein associated	lettuce big-vein associated virus
	varicosavirus*	$(LBVaV)^{14}$
Vesiculovirus	Alagoas vesiculovirus	vesicular stomatitis Alagoas virus (VSAV)
	Carajas vesiculovirus	Carajás virus (CJSV)
	Chandipura vesiculovirus	Chandipura virus (CHPV)
	Cocal vesiculovirus	Cocal virus (COCV)

	Indiana vesiculovirus*	vesicular stomatitis Indiana virus (VSIV)
	Isfahan vesiculovirus	Isfahan virus (ISFV)
	Maraba vesiculovirus	Maraba virus (MARAV)
	New Jersey vesiculovirus	vesicular stomatitis New Jersey virus
		(VSNJV)
	Piry vesiculovirus	Piry virus (OIRYV)
Unassigned	Flanders virus	Flanders virus (FLAV)
	Ngaingan virus	Ngaingan virus (NGAV)
	Wongabel virus	Wongabel virus (WONV)
	Family Sunviridae	
Sunshinevirus	Reptile sunshinevirus 1*	Sunshine Coast virus (SunCV)
	Unassigned	
Anphevirus	Xinchng anphevirus*	Xīnchéng mosquito virus (XcMV)
Arlivirus	Lishi arlivirus*	Líshí spider virus 2 (LsSV-2)
Chengtivirus	Tacheng chengtivirus*	Tăchéng tick virus 6 (TcTV-6)

	Crustavirus	Wenzhou crustavirus*	Wēnzhōu crab virus 1 (WzCV-1)
	Wastrivirus	Sanxia wastrivirus*	Sānxiá water strider virus 4 (SxWSV-4)
1	<sup>1</sup> Includes: Newcastle disease virus (NDV) and pigeon paramyxovirus; <sup>2</sup> synonym: anaconda paramyxovirus; <sup>3</sup> synonym: GH-M74a		nda paramyxovirus; <sup>3</sup> synonym: GH-M74a
2	virus; <sup>4</sup> abbreviation as recently introduced in [26]; <sup>5</sup> historically, an isolate from a samango monkey (Cercopithecus mitis) in 1963 was		
3	long classified as a distinct species called simian agent 10 (SA-10), but was shown later to be HPIV-3. SA-10 was sometimes called		
4	SV-10; <sup>6</sup> synonym: murine parainfluenza virus 1; <sup>7</sup> synonym: bat paramyxovirus; <sup>8</sup> synonym: simian virus 5; <sup>9</sup> synonym: porcine		
5	rubulavirus; <sup>10</sup> synonyms: avian pneumovirus, turkey rhinotracheitis virus; <sup>11</sup> synonyms: citrus leprosis virus nuclear type, citrus		
6	necrotic spot virus; <sup>12</sup> synonym Tzipori virus	s; <sup>13</sup> synonyms: Egtved virus, Paralichthys oliv	vaceus rhabdovirus; <sup>14</sup> synonym: tobacco stunt
7	virus.		

## 1 COMPLIANCE WITH ETHICAL STANDARDS

The views and conclusions contained in this document are those of the authors and should not be 2 interpreted as necessarily representing the official policies, either expressed or implied, of the US 3 Department of the Army, the US Department of Defense, the US Department of Health and 4 Human Services, the Department of Homeland Security (DHS) Science and Technology 5 Directorate (S&T) or of the institutions and companies affiliated with the authors. In no event 6 shall any of these entities have any responsibility or liability for any use, misuse, inability to use, 7 or reliance upon the information contained herein. The US departments do not endorse any 8 9 products or commercial services mentioned in this publication. Funding 10 This work was supported in part through Battelle Memorial Institute's prime contract with the 11 US National Institute of Allergy and Infectious Diseases (NIAID) under Contract No. 12 HHSN272200700016I. A subcontractor to Battelle Memorial Institute who performed this work 13 is: J.H.K., an employee of Tunnell Government Services, Inc. This work was also funded in part 14 under Contract No. HSHQDC-07-C-00020 awarded by DHS S&T for the management and 15 operation of the National Biodefense Analysis and Countermeasures Center (NBACC), a 16 17 Federally Funded Research and Development Center (V.W.-J.); and National Institutes of Health (NIH) contract HHSN272201000040I/HHSN27200004/D04 (N.V., R.B.T.). Y.B. was supported 18 by the Intramural Research Program of the NIH, National Library of Medicine. 19 20 **Conflict of Interest** The authors have no conflicts of interest. 21

22 Ethical approval

1	This article does not contain any studies with human participants or animals performed by any of
2	the authors.

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# 4 **REFERENCES**

5	1.	Adams MJ, Lefkowitz EJ, King AM, Carstens EB (2014) Ratification vote on taxonomic
6		proposals to the International Committee on Taxonomy of Viruses (2014). Arch Virol
7		159:2831-2841

- 8 2. Adams MJ, Lefkowitz EJ, King AM, Bamford DH, Breitbart M, Davison AJ, Ghabrial
- 9 SA, Gorbalenya AE, Knowles NJ, Krell P, Lavigne R, Prangishvili D, Sanfacon H,
- 10 Siddell SG, Simmonds P, Carstens EB (2015) Ratification vote on taxonomic proposals
- to the International Committee on Taxonomy of Viruses (2015). Arch Virol 160:18371850
- 13 3. Bejerman N, Giolitti F, de Breuil S, Trucco V, Nome C, Lenardon S, Dietzgen RG
- 14 (2015) Complete genome sequence and integrated protein localization and interaction
- map for alfalfa dwarf virus, which combines properties of both cytoplasmic and nuclear
   plant rhabdoviruses. Virology 483:275-283
- 17 4. Bishop DHL, Pringle CR (1995) Order *Mononegavirales*. In: Murphy FA, Fauquet CM,
- 18 Bishop DHL, Ghabrial SA, Jarvis AW, Martelli GP, Mayo MA, Summers MD (eds)
- 19 Virus Taxonomy—Sixth Report of the International Committee on Taxonomy of
- 20 Viruses/Archives of Virology Supplement 10. Springer-Verlag, Vienna, Austria, pp 265-

21 267

1	5.	Briand FX, Henry A, Massin P, Jestin V (2012) Complete genome sequence of a novel
2		avian paramyxovirus. J Virol 86:7710
3	6.	Dietzgen RG, Kuhn JH, Clawson AN, Freitas-Astúa J, Goodin MM, Kitajima EW,
4		Kondo H, Wetzel T, Whitfield AE (2014) Dichorhavirus: a proposed new genus for
5		Brevipalpus mite-transmitted, nuclear, bacilliform, bipartite, negative-strand RNA plant
6		viruses. Arch Virol 159:607-619
7	7.	Drexler JF, Corman VM, Müller MA, Maganga GD, Vallo P, Binger T, Gloza-Rausch F,
8		Cottontail VM, Rasche A, Yordanov S, Seebens A, Knörnschild M, Oppong S, Adu
9		Sarkodie Y, Pongombo C, Lukashev AN, Schmidt-Chanasit J, Stöcker A, Borges
10		Carneiro AJ, Erbar S, Maisner A, Fronhoffs F, Buettner R, Kalko EKV, Kruppa T,
11		Franke CR, Kallies R, Yandoko ERN, Herrler G, Reusken C, Hassanin A, Krüger DH,
12		Matthee S, Ulrich RG, Leroy EM, Drosten C (2012) Bats host major mammalian
13		paramyxoviruses. Nat Commun 3:796
14	8.	Easton AJ, Pringle CR (2011) Order Mononegavirales. In: King AMQ, Adams MJ,
15		Carstens EB, Lefkowitz EJ (eds) Virus Taxonomy-Ninth Report of the International
16		Committee on Taxonomy of Viruses. Elsevier/Academic Press, London, United
17		Kingdom, pp 653-657
18	9.	Guo J, Tizard I (2015) The genome sequence of parrot bornavirus 5. Virus Genes 51:430-
19		433
20	10.	Hyndman TH, Marschang RE, Wellehan JF, Jr., Nicholls PK (2012) Isolation and
21		molecular identification of Sunshine virus, a novel paramyxovirus found in Australian
22		snakes. Infect Genet Evol 12:1436-1446

1	11.	Kuhn JH, Bekal S, Caì Y, Clawson AN, Domier LL, Herrel M, Jahrling PB, Kondo H,
2		Lambert KN, Mihindukulasuriya KA, Nowotny N, Radoshitzky SR, Schneider U,
3		Staeheli P, Suzuki N, Tesh RB, Wang D, Wang L-F, Dietzgen RG (2013) Nyamiviridae:
4		proposal for a new family in the order Mononegavirales. Arch Virol 158:2209–2226
5	12.	Kuhn JH, Dürrwald R, Bào Y, Briese T, Carbone K, Clawson AN, deRisi JL, Garten W,
6		Jahrling PB, Kolodziejek J, Rubbenstroth D, Schwemmle M, Stenglein M, Tomonaga K,
7		Weissenböck H, Nowotny N (2015) Taxonomic reorganization of the family
8		Bornaviridae. Arch Virol 160:621-632
9	13.	Kumar S, Collins PL, Samal SK (2010) Identification of simian agent 10 as human
10		parainfluenza virus type 3 suggests transmission of a human virus to an African monkey.
11		J Virol 84:13068-13070
12	14.	Lau SK, Woo PC, Wu Y, Wong AY, Wong BH, Lau CC, Fan RY, Cai JP, Tsoi HW,
13		Chan KH, Yuen KY (2013) Identification and characterization of a novel paramyxovirus,
14		porcine parainfluenza virus 1, from deceased pigs. J Gen Virol 94:2184-2190
15	15.	Li CX, Shi M, Tian JH, Lin XD, Kang YJ, Chen LJ, Qin XC, Xu J, Holmes EC, Zhang
16		YZ (2015) Unprecedented genomic diversity of RNA viruses in arthropods reveals the
17		ancestry of negative-sense RNA viruses. Elife 4:e05378
18	16.	Liu L, Xie J, Cheng J, Fu Y, Li G, Yi X, Jiang D (2014) Fungal negative-stranded RNA
19		virus that is related to bornaviruses and nyaviruses. Proc Natl Acad Sci U S A
20		111:12205-12210
21	17.	Marsh GA, de Jong C, Barr JA, Tachedjian M, Smith C, Middleton D, Yu M, Todd S,
22		Foord AJ, Haring V, Payne J, Robinson R, Broz I, Crameri G, Field HE, Wang LF (2012)
23		Cedar virus: a novel henipavirus isolated from Australian bats. PLoS Pathog 8:e1002836
		Mononegavirales Taxonomy Update 27

1	18.	Marton S, Bányai K, Gál J, Ihász K, Kugler R, Lengyel G, Jakab F, Bakonyi T, Farkas
2		SL (2015) Coding-complete sequencing classifies parrot bornavirus 5 into a novel virus
3		species. Arch Virol 160:2763-2768
4	19.	Miller PJ, Afonso CL, Spackman E, Scott MA, Pedersen JC, Senne DA, Brown JD,
5		Fuller CM, Uhart MM, Karesh WB, Brown IH, Alexander DJ, Swayne DE (2010)
6		Evidence for a new avian paramyxovirus serotype 10 detected in rockhopper penguins
7		from the Falkland Islands. J Virol 84:11496-11504
8	20.	Pringle CR (1991) Order Mononegavirales. In: Francki RIB, Fauquet CM, Knudson DL,
9		Brown F (eds) Classification and Nomenclature of Viruses—Fifth Report of the
10		International Committee on Taxonomy of Viruses/Archives of Virology Supplementum
11		2. Springer-Verlag, Vienna, Austria, pp 239-241
12	21.	Pringle CR, Alexander DJ, Billeter MA, Collins PL, Kingsbury DW, Lipkind MA, Nagai
13		Y, Orvell C, Rima B, Rott R, ter Meulen V (1991) The order Mononegavirales. Arch
14		Virol 117:137-140
15	22.	Pringle CR (1996) Virus taxonomy 1996—a bulletin from the Xth International Congress
16		of Virology in Jerusalem. Arch Virol 141:2251-2256
17	23.	Pringle CR (1997) The order Mononegavirales—current status. Arch Virol 142:2321-
18		2326
19	24.	Pringle CR (2000) Order Mononegavirales. In: van Regenmortel MHV, Fauquet CM,
20		Bishop DHL, Carstens EB, Estes MK, Lemon SM, Maniloff J, Mayo MA, McGeoch DJ,
21		Pringle CR, Wickner RB (eds) Virus Taxonomy—Seventh Report of the International
22		Committee on Taxonomy of Viruses. Academic Press, San Diego, California, USA, pp
23		525-530

25.	Pringle CR (2005) Order Mononegavirales. In: Fauquet CM, Mayo MA, Maniloff J,
	Desselberger U, Ball LA (eds) Virus Taxonomy-Eighth Report of the International
	Committee on Taxonomy of Viruses. Elsevier/Academic Press, San Diego, California,
	USA, pp 609-614
26.	Sharp CR, Nambulli S, Acciardo AS, Rennick LJ, Drexler JF, Rima BK, Williams T,
	Duprex WP (2016) Chronic infection of domestic cats with feline morbillivirus, United
	States. Emerg Infect Dis 22:760-762
27.	Stenglein MD, Leavitt EB, Abramovitch MA, McGuire JA, DeRisi JL (2014) Genome
	sequence of a bornavirus recovered from an African garter snake (Elapsoidea loveridgei).
	Genome Announc 2:e00779-00714
28.	Terregino C, Aldous EW, Heidari A, Fuller CM, De Nardi R, Manvell RJ, Beato MS,
	Shell WM, Monne I, Brown IH, Alexander DJ, Capua I (2013) Antigenic and genetic
	analyses of isolate APMV/wigeon/Italy/3920-1/2005 indicate that it represents a new
	avian paramyxovirus (APMV-12). Arch Virol 158:2233-2243
29.	Van Regenmortel MH, Burke DS, Calisher CH, Dietzgen RG, Fauquet CM, Ghabrial SA,
	Jahrling PB, Johnson KM, Holbrook MR, Horzinek MC, Keil GM, Kuhn JH, Mahy BW,
	Martelli GP, Pringle C, Rybicki EP, Skern T, Tesh RB, Wahl-Jensen V, Walker PJ,
	Weaver SC (2010) A proposal to change existing virus species names to non-Latinized
	binomials. Arch Virol 155:1909-1919
30.	Woo PC, Lau SK, Wong BH, Fan RY, Wong AY, Zhang AJ, Wu Y, Choi GK, Li KS,
	Hui J, Wang M, Zheng BJ, Chan KH, Yuen KY (2012) Feline morbillivirus, a previously
	undescribed paramyxovirus associated with tubulointerstitial nephritis in domestic cats.
	Proc Natl Acad Sci U S A 109:5435-5440
	<ul> <li>26.</li> <li>27.</li> <li>28.</li> <li>29.</li> </ul>

- 1 31. Wu Z, Yang L, Yang F, Ren X, Jiang J, Dong J, Sun L, Zhu Y, Zhou H, Jin Q (2014)
- 2 Novel henipa-like virus, Mojiang paramyxovirus, in rats, China, 2012. Emerg Infect Dis
- 3 20:1064-1066