



Taxonomy of the order *Mononegavirales*: update 2017

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Received: 19 February 2017 / Accepted: 26 February 2017 / Published online: 7 April 2017

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Abstract In 2017, the order *Mononegavirales* was expanded by the inclusion of a total of 69 novel species. Five new rhabdovirus genera and one new nyamivivirus genus were established to harbor 41 of these species, whereas the remaining new species were assigned to already established genera. Furthermore, non-Latinized

binomial species names replaced all paramyxovirus and pneumovirus species names, thereby accomplishing application of binomial species names throughout the entire order. This article presents the updated taxonomy of the order *Mononegavirales* as now accepted by the International Committee on Taxonomy of Viruses (ICTV).

Members of the International Committee on Taxonomy of Viruses (ICTV) *Bornnaviridae*, *Filoviridae*, *Mononegavirales*, *Nyamiviridae*, *Paramyxoviridae*, and *Rhabdoviridae* Study Groups who co-authored this manuscript are listed in “Acknowledgements”.

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Introduction

The virus order *Mononegavirales* was established in 1991 to accommodate related viruses with nonsegmented, linear, single-stranded negative-sense RNA genomes. These viruses were initially assigned to three mononegavirus families: *Filoviridae*, *Paramyxoviridae*, and *Rhabdoviridae* [32, 33]. The families *Bornaviridae* and *Nyamiviridae* joined the order in 1996 [34] and 2014 [1, 25], respectively, followed by the families *Mymonaviridae*, *Pneumoviridae*, and *Sunviridae* in 2016 [2]. The order was continuously amended in 1995 [7], 1997 [35], 2000 [36], 2005 [37], 2011 [17], and 2016 [2]. In 2016, the Study Groups of the International Committee on Taxonomy of Viruses (ICTV) responsible for the taxonomy of the order and its eight families assigned unclassified mononegaviruses to existing or novel taxa and continued efforts to streamline order nomenclature. Here we present the changes that were (re)proposed via official ICTV taxonomic proposals (TaxoProps) at <http://www.ictvonline.org/> in 2016 and that were accepted by the ICTV Executive Committee. These changes are official ICTV taxonomy as of 2017.

Taxonomic changes at the order level

No changes were made at the order level.

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Taxonomic changes at the family level

Bornaviridae

The family *Bornaviridae* remains monogeneric but was expanded in 2017 by one species (*Mammalian 2 bornavirus*) for the newly discovered variegated squirrel bornavirus 1 [23] (TaxoProp 2016.013aM.A.v1.Bornavirus_sp).

Filoviridae

No changes were made to this family.

Mymonaviridae

No changes were made to this family.

Nyamiviridae

The family *Nyamiviridae* was expanded to include a third genus (*Peropivirus*) including the new species *Pteromalus puparum peropivirus* for a virus recently discovered in parasitoid wasps, *Pteromalus puparum* negative-strand RNA virus 1 [45] (TaxoProp 2016.015a-dM.A.v1.Peropivirus).

Paramyxoviridae

The non-Latinized binomial species name format [40] was applied throughout the family *Paramyxoviridae* (TaxoProp

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2016.011aM.A.v2.Paramyxoviridae_spren). In addition, the genus *Avulavirus* was expanded by one novel species, *Avian avulavirus 13*, for avian paramyxovirus 13 discovered in geese [19, 24, 47] (TaxoProp 2016.001a,bM.A.-v2.Avulavirus_spren). The genus *Rubulavirus* was expanded by 10 species to accommodate two long-known viruses (e.g., Menangle virus, Tioman virus) and eight newly discovered bat viruses (bat mumps virus, Achimota viruses 1 and 2, Sosuga virus, Teviot virus, Tuhoko viruses 1–3) [3, 5, 11, 12, 16, 26] (TaxoProp 2015.016aM.A.v3.-Rubulavirus_10sp). Finally, the previous species name *Newcastle disease virus* was changed to *Avian avulavirus 1* to reflect the fact that “Newcastle disease virus” is but one strain of the member virus of this species, avian paramyxovirus 1 (APMV-1) [46] (TaxoProp 2016.001a,bM.A.v2.Avulavirus_spren).

Pneumoviridae

The non-Latinized binomial species name format [40] was applied throughout the family *Pneumoviridae* (TaxoProp 2016.012aM.A.v1.Pneumoviridae_spren).

Rhabdoviridae

The most extensive changes were introduced into the family *Rhabdoviridae*. The new genus *Almendravirus* was created to include five new species (TaxoProp 2016.002a-dM.A.v3.Almendravirus) to accommodate mosquito viruses discovered from 2009–2013 (Arboretum virus, Balsa virus, Coot Bay virus, Puerto Almendras virus, and Rio Chico virus) [13, 43, 44]. The new genus *Curiovirus*, including four new species, was created for unclassified dipteran viruses known since the 1970s–80s (Curionopolis virus, Iriri virus, Itacaiunas virus, Rochambeau virus) (TaxoProp 2016.003a-dM.A.v3.Curiovirus) [14, 15, 44]. Twelve mostly dipteran viruses (Hart Park virus, Gray Lodge virus, Joinjakaka virus, Kameise virus, La Joya virus, Landjia virus, Manitoba virus, Marco virus, Mosqueiro virus, Mossuril virus, Ord River virus, and Parry Creek virus), isolated in the 1950s to late 1970s, were classified into twelve new species in the new genus *Hapavirus*. The previously free-floating rhabdovirus species *Flanders virus*, *Ngaingan virus*, and *Wongabel virus* were moved into the genus *Hapavirus* and renamed accordingly (*Flanders hapavirus*, *Ngaingan hapavirus*, and *Wongabel*

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hapavirus, respectively) [4, 21, 44] (TaxoProp 2016.005a-dM.A.v4.Hapavirus). The largest newly created rhabdovirus genus is *Ledantivirus*. This genus includes 14 new species for Barur virus, Fikirini virus, Fukuoka virus, Kern Canyon virus, Keuraliba virus, Kolente virus, Kumasi rhabdovirus, Le Dantec virus, Mount Elgon bat virus, Nkolbisson virus, Nishimuro virus, Oita virus, Wūhàn louse fly virus 5, and Yōngjiā tick virus 2, which were isolated from or detected in arthropods and/or mammals [6, 10, 18, 28, 38, 44] (TaxoProp 2016.006a-dM.A.v2.Ledantivirus). Finally, the novel genus *Sripuvirus* was created to include five new species for sandfly and lizard viruses (Almpiwar virus, Chaco virus, Niakha virus, Sena Madureira virus, and Sripur virus) [29, 41, 44] (TaxoProp 2016.007a-dM.A.v5.Sripuvirus).

The existing genus *Cytorhabdovirus* was expanded by inclusion of the novel species *Colocasia bobone disease-associated cytorhabdovirus* for a plant virus first described in 1973 in the taro plant (*Colocasia esculenta*) [22] (TaxoProp 2016.017aM.A.v1.Cytorhabdovirus_sp). The genus *Ephemerovirus* was enlarged by three species to accommodate the long-known bovid viruses Kimberley and Koolpinyah, and the mosquito-borne Malakal and Yata viruses [8, 9] (TaxoProp 2016.004aM.A.v2.Ephemerovirus_3sp). The recently described, possibly human, Bas-Congo virus and Ekpoma viruses 1 and 2, and the biting midge-borne Sweetwater Branch virus were classified into four new species of the genus

Tibrovirus [20, 27, 39, 44] (TaxoProp 2016.008aM.A.v3.Tibrovirus_4sp). Klamath virus, discovered in 1962 in a vole, was assigned to a novel species in the genus *Tupavivirus* [44] (TaxoProp 2016.009aM.A.v2.Tupavivirus_sp), and seven new species were added to the genus *Vesiculovirus* for American bat vesiculovirus (isolated from bats in 2008) and Jurona virus, Malpais Spring virus, Morreton virus, Perinet virus, Radi virus, and Yug Bogdanovac virus (isolated from mosquitoes or sandflies in 1962–1986) [22, 30, 31, 42, 44] (TaxoProp 016.010aM.A.v3.Vesiculovirus_7sp).

Finally, the existing species names *Oncorhynchus 1 novirhabdovirus* (formerly *Infectious hematopoietic necrosis virus*) and *Oncorhynchus 2 novirhabdovirus* (formerly *Viral hemorrhagic septicemia virus*) were changed to *Salmonid novirhabdovirus* and *Piscine novirhabdovirus*, respectively, to be more reflective of the host spectrum of their members. (2016.018aM.A.v1.Novirhabdovirus_spren).

Sunviridae

In 2017, no changes were made to this family.

Summary

A summary of the current, ICTV-accepted taxonomy of the order *Mononegavirales* is presented in Table 1.

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Table 1 ICTV-accepted taxonomy of the order *Mononegavirales* as of 2017

Genus	Species [*]	Virus (Abbreviation) [†]
Family Bornaviridae		
<i>Bornavirus</i>	<i>Elapid 1 bornavirus</i>	Loveridge's garter snake virus 1 (LGSV-1)
	<i>Mammalian 1 bornavirus*</i>	Borna disease virus 1 (BoDV-1)
	<i>Mammalian 2 bornavirus</i>	Borna disease virus 2 (BoDV-2)
	<i>Passeriform 1 bornavirus</i>	variegated squirrel bornavirus 1 (VSBV-1)
	<i>Passeriform 2 bornavirus</i>	canary bornavirus 1 (CnBV-1)
	<i>Psittaciform 1 bornavirus</i>	canary bornavirus 2 (CnBV-2)
	<i>Psittaciform 2 bornavirus</i>	canary bornavirus 3 (CnBV-3)
	<i>Waterbird 1 bornavirus</i>	estrildid finch bornavirus 1 (EsBV-1)
		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)
		parrot bornavirus 5 (PaBV-5)
		aquatic bird bornavirus 1 (ABBV-1)
		aquatic bird bornavirus 2 (ABBV-2)
Family Filoviridae		
<i>Cuevavirus</i>	<i>Lloviu cuevavirus*</i>	Lloviu virus (LLOV)
<i>Ebolavirus</i>	<i>Bundibugyo ebolavirus</i>	Bundibugyo virus (BDBV)
	<i>Reston ebolavirus</i>	Reston virus (RESTV)
	<i>Sudan ebolavirus</i>	Sudan virus (SUDV)
	<i>Tai Forest ebolavirus</i>	Tai Forest virus (TAFV)
	<i>Zaire ebolavirus*</i>	Ebola virus (EBOV)
<i>Marburgvirus</i>	<i>Marburg marburgvirus*</i>	Marburg virus (MARV)
		Ravn virus (RAVV)
Family Mymonaviridae		
	<i>Sclerotimonavirus</i>	Sclerotinia sclerotiorum negative-stranded RNA virus 1 (SsNSRV-1)
Family Nyamiviridae		
<i>Nyavirus</i>	<i>Midway nyavirus</i>	Midway virus (MIDWV)
	<i>Nyamanini nyavirus*</i>	Nyamanini virus (NYMV)
	<i>Sierra Nevada nyavirus</i>	Sierra Nevada virus (SNVV)
<i>Peropivirus</i>	<i>Pteromalus puparum peropivirus*</i>	Pteromalus puparum negative-strand RNA virus 1 (PpNSRV-1)
<i>Socyvirus</i>	<i>Soybean cyst nematode socyvirus*</i>	soybean cyst nematode virus 1 (SbCNV-1)
Family Paramyxoviridae		
<i>Aquaparamyxovirus</i>	<i>Salmon aquaparamyxovirus*</i>	Atlantic salmon paramyxovirus (AsaPV)
<i>Avulavirus</i>	<i>Avian avulavirus 1*</i>	avian paramyxovirus 1 (APMV-1) ¹
	<i>Avian avulavirus 2</i>	avian paramyxovirus 2 (APMV-2)
	<i>Avian avulavirus 3</i>	avian paramyxovirus 3 (APMV-3)
	<i>Avian avulavirus 4</i>	avian paramyxovirus 4 (APMV-4)
	<i>Avian avulavirus 5</i>	avian paramyxovirus 5 (APMV-5)
	<i>Avian avulavirus 6</i>	avian paramyxovirus 6 (APMV-6)
	<i>Avian avulavirus 7</i>	avian paramyxovirus 7 (APMV-7)
	<i>Avian avulavirus 8</i>	avian paramyxovirus 8 (APMV-8)
	<i>Avian avulavirus 9</i>	avian paramyxovirus 9 (APMV-9)
	<i>Avian avulavirus 10</i>	avian paramyxovirus 10 (APMV-10)
	<i>Avian avulavirus 11</i>	avian paramyxovirus 11 (APMV-11)
	<i>Avian avulavirus 12</i>	avian paramyxovirus 12 (APMV-12)
	<i>Avian avulavirus 13</i>	avian paramyxovirus 13 (APMV-13)
<i>Ferlavirius</i>	<i>Reptilian ferlavirius*</i>	Fer-de-Lance virus (FDLV) ²
<i>Henipavirus</i>	<i>Cedar henipavirus</i>	Cedar virus (CedV)

Table 1 continued

Genus	Species [¶]	Virus (Abbreviation) [¶]
	<i>Ghanaian bat henipavirus</i>	Kumasi virus (KV) ³
	<i>Hendra henipavirus*</i>	Hendra virus (HeV)
	<i>Mojiang henipavirus</i>	Mòjiāng virus (MojV)
	<i>Nipah henipavirus</i>	Nipah virus (NiV)
<i>Morbillivirus</i>	<i>Canine morbillivirus</i>	canine distemper virus (CDV)
	<i>Cetacean morbillivirus</i>	cetacean morbillivirus (CeMV)
	<i>Feline morbillivirus</i>	feline morbillivirus (FeMV)
	<i>Measles morbillivirus*</i>	measles virus (MeV)
	<i>Small ruminant morbillivirus</i>	peste-des-petits-ruminants virus (PPRV)
	<i>Phocine morbillivirus</i>	phocine distemper virus (PDV)
	<i>Rinderpest morbillivirus</i>	rinderpest virus (RPV)
<i>Respirovirus</i>	<i>Bovine respirovirus 3</i>	bovine parainfluenza virus 3 (BPIV-3)
	<i>Human respirovirus 1</i>	human parainfluenza virus 1 (HPIV-1)
	<i>Human respirovirus 3</i>	human parainfluenza virus 3 (HPIV-3)
	<i>Porcine respirovirus 1</i>	porcine parainfluenza virus 1 (PPIV-1)
	<i>Murine respirovirus*</i>	Sendai virus (SeV) ⁴
<i>Rubulavirus</i>	<i>Achimota rubulavirus 1</i>	Achimota virus 1 (AchPV-1)
	<i>Achimota rubulavirus 2</i>	Achimota virus 2 (AchPV-2)
	<i>Bat mumps rubulavirus</i>	bat mumps virus (BMV) ⁵
	<i>Canine rubulavirus</i>	parainfluenza virus 5 (PIV-5) ⁶
	<i>Human rubulavirus 2</i>	human parainfluenza virus 2 (HPIV-2)
	<i>Human rubulavirus 4</i>	human parainfluenza virus 4a (HPIV-4a)
	<i>Mapuera rubulavirus</i>	human parainfluenza virus 4b (HPIV-4b)
	<i>Menangle rubulavirus</i>	Mapuera virus (MapV)
	<i>Mumps rubulavirus*</i>	Menangle virus (MenPV)
	<i>Porcine rubulavirus</i>	mumps virus (MuV)
	<i>Simian rubulavirus</i>	La Piedad Michoacán Mexico virus (LPMV) ⁷
	<i>Sosuga rubulavirus</i>	simian virus 41 (SV-41)
	<i>Teviot rubulavirus</i>	Sosuga virus
	<i>Tioman rubulavirus</i>	Teviot virus (TevPV)
	<i>Tuhoko rubulavirus 1</i>	Tioman virus (TioPV)
	<i>Tuhoko rubulavirus 2</i>	Tuhoko virus 1 (ThkPV-1)
	<i>Tuhoko rubulavirus 3</i>	Tuhoko virus 2 (ThkPV-2)
		Tuhoko virus 3 (ThkPV-3)
Family <i>Pneumoviridae</i>		
<i>Metapneumovirus</i>	<i>Avian metapneumovirus*</i>	avian metapneumovirus (AMPV) ⁸
	<i>Human metapneumovirus</i>	human metapneumovirus (HMPV)
<i>Orthopneumovirus</i>	<i>Bovine orthopneumovirus</i>	bovine respiratory syncytial virus (BRSV)
	<i>Human orthopneumovirus*</i>	human respiratory syncytial virus A2 (HRSV-A2)
	<i>Murine orthopneumovirus</i>	human respiratory syncytial virus B1 (HRSV-B1)
		murine pneumonia virus (MPV)
Family <i>Rhabdoviridae</i>		
<i>Almendravirus</i>	<i>Arboretum almendravirus</i>	Arboretum virus (ABTV)
	<i>Balsa almendravirus</i>	Balsa virus (BALV)
	<i>Coot Bay almendravirus</i>	Coot Bay virus (CBV)
	<i>Puerto Almendras almendravirus*</i>	Puerto Almendras virus (PTAMV)
	<i>Rio Chico almendravirus</i>	Rio Chico virus (RCHV)
<i>Curiovirus</i>	<i>Curionopolis curiovirus*</i>	Curionopolis virus (CURV)
	<i>Iriri curiovirus</i>	Iriri virus (IRIRV)
	<i>Itacaiunas curiovirus</i>	Itacaiunas virus (ITAV)
	<i>Rochambeau curiovirus</i>	Rochambeau virus (RBUV)
<i>Cytorhabdovirus</i>	<i>Alfalfa dwarf cytorhabdovirus</i>	alfalfa dwarf virus (ADV)

Table 1 continued

Genus	Species ^a	Virus (Abbreviation) ^b
	<i>Barley yellow striate mosaic cytorhabdovirus</i>	barley yellow striate mosaic virus (BYSMV)
	<i>Broccoli necrotic yellows cytorhabdovirus</i>	broccoli necrotic yellows virus (BNYV)
	<i>Colocasia bobone disease-associated cytorhabdovirus</i>	Colocasia bobone disease-associated virus (CBDAV)
	<i>Festuca leaf streak cytorhabdovirus</i>	festuca leaf streak virus (FLSV)
	<i>Lettuce necrotic yellows cytorhabdovirus*</i>	lettuce necrotic yellows virus (LNYV)
	<i>Lettuce yellow mottle cytorhabdovirus</i>	lettuce yellow mottle virus (LYMoV)
	<i>Northern cereal mosaic cytorhabdovirus</i>	northern cereal mosaic virus (NCMV)
	<i>Sonchus cytorhabdovirus 1</i>	sonchus virus (SonV)
	<i>Strawberry crinkle cytorhabdovirus</i>	strawberry crinkle virus (SCV)
	<i>Wheat American striate mosaic cytorhabdovirus</i>	wheat American striate mosaic virus (WASMV)
<i>Dichorhavirus</i>	<i>Coffee ringspot dichorhavirus</i>	coffee ringspot virus (CoRSV)
	<i>Orchid fleck dichorhavirus*</i>	orchid fleck virus (OFV) ⁹
<i>Ephemerovirus</i>	<i>Adelaide River ephemerovirus</i>	Adelaide River virus (ARV)
	<i>Berrimah ephemerovirus</i>	Berrimah virus (BRMV)
	<i>Bovine fever ephemerovirus*</i>	bovine ephemeral fever virus (BEFV) ¹⁰
	<i>Kimberley ephemerovirus</i>	Kimberley virus (KIMV)
	<i>Koolpinyah ephemerovirus</i>	Koolpinyah virus (KOOLV)
	<i>Kotonkan ephemerovirus</i>	kotonkan virus (KOTV)
	<i>Obodhiang ephemerovirus</i>	Obodhiang virus (OBOV)
	<i>Yata ephemerovirus</i>	Yata virus (YATV)
<i>Hapavirus</i>	<i>Flanders hapavirus</i>	Flanders virus (FLAV)
	<i>Hart Park hapavirus</i>	Hart Park virus (HPV)
	<i>Gray Lodge hapavirus</i>	Gray Lodge virus (GLOV)
	<i>Joinjakaka hapavirus</i>	Joinjakaka virus (JOIV)
	<i>La Joya hapavirus</i>	La Joya virus (LJV)
	<i>Kamese hapavirus</i>	Kamese virus (KAMV)
	<i>Landjia hapavirus</i>	Landjia virus (LANV = LJAV)
	<i>Manitoba hapavirus</i>	Manitoba virus (MANV = MNTBV)
	<i>Marco hapavirus</i>	Marco virus (MCOV)
	<i>Mosqueiro hapavirus</i>	Mosqueiro virus (MQOV)
	<i>Mossuril hapavirus</i>	Mossuril virus (MOSV)
	<i>Ngaingan hapavirus</i>	Ngaingan virus (NGAV)
	<i>Ord River hapavirus</i>	Ord River virus (ORV)
	<i>Parry Creek hapavirus</i>	Parry Creek virus (PCV)
	<i>Wongabel hapavirus*</i>	Wongabel virus (WONV)
<i>Ledantivirus</i>	<i>Barur ledantivirus</i>	Barur virus (BARV)
	<i>Fikirini ledantivirus</i>	Fikirini virus (FKRV)
	<i>Fukuoka ledantivirus</i>	Fukuoka virus (FUKV)
	<i>Kern Canyon ledantivirus</i>	Kern Canyon virus (KCV)
	<i>Keuraliba ledantivirus</i>	Keuraliba virus (KEUV)
	<i>Kolente ledantivirus</i>	Kolente virus (KOLEV)
	<i>Kumasi ledantivirus</i>	Kumasi rhabdovirus (KRV)
	<i>Le Dantec ledantivirus*</i>	Le Dantec virus (LDV)
	<i>Mount Elgon bat ledantivirus</i>	Mount Elgon bat virus (MEBV)
	<i>Nkolbisson ledantivirus</i>	Nkolbisson virus (NKOV)
	<i>Nishimuro ledantivirus</i>	Nishimuro virus (NISV) ¹¹
	<i>Oita ledantivirus</i>	Oita virus (OITAV)
	<i>Wuhan ledantivirus</i>	Wūhān louse fly virus 5 (WLFV-5)
	<i>Yongjia ledantivirus</i>	Yōngjiā tick virus 2 (YTV-2)
<i>Lyssavirus</i>	<i>Aravan lyssavirus</i>	Aravan virus (ARAV)
	<i>Australian bat lyssavirus</i>	Australian bat lyssavirus (ABLV)
	<i>Bokeloh bat lyssavirus</i>	Bokeloh bat lyssavirus (BBLV)

Table 1 continued

Genus	Species [*]	Virus (Abbreviation) [†]
	<i>Duvenhage lyssavirus</i>	Duvenhage virus (DUVV)
	<i>European bat 1 lyssavirus</i>	European bat lyssavirus 1 (EBLV-1)
	<i>European bat 2 lyssavirus</i>	European bat lyssavirus 2 (EBLV-2)
	<i>Ikoma lyssavirus</i>	Ikoma lyssavirus (IKOV)
	<i>Irkut lyssavirus</i>	Irkut virus (IRKV)
	<i>Khujand lyssavirus</i>	Khujand virus (KHUV)
	<i>Lagos bat lyssavirus</i>	Lagos bat virus (LBV)
	<i>Mokola lyssavirus</i>	Mokola virus (MOKV)
	<i>Rabies lyssavirus*</i>	rabies virus (RABV)
	<i>Shimoni bat lyssavirus</i>	Shimoni bat virus (SHIBV)
	<i>West Caucasian bat lyssavirus</i>	West Caucasian bat virus (WCBV)
<i>Novirhabdovirus</i>	<i>Hirame novirhabdovirus</i>	Hirame rhabdovirus (HIRV)
	<i>Piscine novirhabdovirus</i>	viral hemorrhagic septicemia virus (VHSV) ¹²
	<i>Salmonid novirhabdovirus*</i>	infectious hematopoietic necrosis virus (IHNV)
	<i>Snakehead novirhabdovirus</i>	snakehead rhabdovirus (SHRV)
<i>Nucleorhabdovirus</i>	<i>Datura yellow vein nucleorhabdovirus</i>	datura yellow vein virus (DYVV)
	<i>Eggplant mottled dwarf nucleorhabdovirus</i>	eggplant mottled dwarf virus (EMDV)
	<i>Maize fine streak nucleorhabdovirus</i>	maize fine streak virus (MSFV)
	<i>Maize Iranian mosaic nucleorhabdovirus</i>	maize Iranian mosaic virus (MIMV)
	<i>Maize mosaic nucleorhabdovirus</i>	maize mosaic virus (MMV)
	<i>Potato yellow dwarf nucleorhabdovirus*</i>	potato yellow dwarf virus (PYDV)
	<i>Rice yellow stunt nucleorhabdovirus</i>	rice yellow stunt virus (RYSV)
	<i>Sonchus yellow net nucleorhabdovirus</i>	rice transitory yellowing virus (RTYV)
	<i>Sowthistle yellow vein nucleorhabdovirus</i>	sonchus yellow net virus (SYNV)
	<i>Taro vein chlorosis nucleorhabdovirus</i>	sowthistle yellow vein virus (SYVV)
<i>Perhabdovirus</i>	<i>Anguillid perhabdovirus</i>	taro vein chlorosis virus (TaVCV)
	<i>Perch perhabdovirus*</i>	eel virus European X (EVEX)
	<i>Sea trout perhabdovirus</i>	perch rhabdovirus (PRV)
<i>Sigmavirus</i>	<i>Drosophila affinis sigmavirus</i>	lake trout rhabdovirus (LTRV)
	<i>Drosophila ananassae sigmavirus</i>	Drosophila affinis sigmavirus (DAffSV)
	<i>Drosophila immigrans sigmavirus</i>	Drosophila ananassae sigmavirus (DAAnSV)
	<i>Drosophila melanogaster sigmavirus*</i>	Drosophila immigrans sigmavirus (DIImmSV)
	<i>Drosophila obscura sigmavirus</i>	Drosophila melanogaster sigmavirus (DMelSV)
	<i>Drosophila tristis sigmavirus</i>	Drosophila obscura sigmavirus (DObsSV)
	<i>Muscina stabulans sigmavirus</i>	Drosophila tristis sigmavirus (DTriSV)
<i>Sprivivirus</i>	<i>Carp sprivivirus*</i>	Muscina stabulans sigmavirus (MStaSV)
	<i>Pike fry sprivivirus</i>	spring viremia of carp virus (SVCV)
		grass carp rhabdovirus (GrCRV)
		pike fry rhabdovirus (PFRV)
		tench rhabdovirus (TenRV)
<i>Sripuvirus</i>	<i>Almpiwar sripuvirus</i>	Almpiwar virus (ALMV)
	<i>Chaco sripuvirus</i>	Chaco virus (CHOV)
	<i>Niakha sripuvirus*</i>	Niakha virus (NIAV)
	<i>Sena Madueira sripuvirus</i>	Sena Madueira virus (SMV)
	<i>Sripur sripuvirus</i>	Sripur virus (SRIV)
<i>Tibrovirus</i>	<i>Bas-Congo tibrovirus</i>	Bas-Congo virus (BASV)
	<i>Coastal Plains tibrovirus</i>	Coastal Plains virus (CPV)
	<i>Ekpoma 1 tibrovirus</i>	Ekpoma virus 1 (EKV-1)
	<i>Ekpoma 2 tibrovirus</i>	Ekpoma virus 2 (EKV-2)
	<i>Sweetwater Branch tibrovirus</i>	Sweetwater Branch virus (SWBV)
	<i>Tibrogargan tibrovirus*</i>	Bivens Arm virus (BAV)
		Tibrogargan virus (TIBV)

Table 1 continued

Genus	Species [*]	Virus (Abbreviation) [†]
<i>Tupavirus</i>	<i>Durham tupavirus</i> *	Durham virus (DURV)
	<i>Klamath tupavirus</i>	Klamath virus (KLAV)
	<i>Tupaia tupavirus</i>	tupaia virus (TUPV)
<i>Varicosavirus</i>	<i>Lettuce big-vein associated varicosavirus</i> *	lettuce big-vein associated virus (LBVaV) ¹³
<i>Vesiculovirus</i>	<i>Alagoas vesiculovirus</i>	vesicular stomatitis Alagoas virus (VSAV)
	<i>American bat vesiculovirus</i>	American bat vesiculovirus (ABVV)
	<i>Carajas vesiculovirus</i>	Carajás virus (CJSV)
	<i>Chandipura vesiculovirus</i>	Chandipura virus (CHPV)
	<i>Cocal vesiculovirus</i>	Cocal virus (COCV)
	<i>Indiana vesiculovirus</i> *	vesicular stomatitis Indiana virus (VSIV)
	<i>Isfahan vesiculovirus</i>	Isfahan virus (ISFV)
	<i>Jurona vesiculovirus</i>	Jurona virus (JURV)
	<i>Malpais Spring vesiculovirus</i>	Malpais Spring virus (MSPV)
	<i>Maraba vesiculovirus</i>	Maraba virus (MARAV)
	<i>Morreton vesiculovirus</i>	Morreton virus (MORV)
	<i>New Jersey vesiculovirus</i>	vesicular stomatitis New Jersey virus (VSNJV)
Unassigned	<i>Perinet vesiculovirus</i>	Perinet virus (PERV)
	<i>Piry vesiculovirus</i>	Piry virus (PIRYV)
	<i>Radi vesiculovirus</i>	Radi virus (RADV)
	<i>Yug Bogdanovac vesiculovirus</i>	Yug Bogdanovac virus (YBV)
	<i>Moussa virus</i>	Moussa virus (MOUV)
Family <i>Sunviridae</i>		
<i>Sunshinevirus</i>	<i>Reptile sunshinevirus 1</i> *	Sunshine Coast virus (SunCV)
Unassigned		
<i>Anphevirus</i>	<i>Xincheng anphevirus</i> *	Xīnchéng mosquito virus (XcMV)
<i>Arlivirus</i>	<i>Lishi arlivirus</i> *	Líshí spider virus 2 (LsSV-2)
<i>Chengtivirus</i>	<i>Tacheng chengtivirus</i> *	Tāchéng tick virus 6 (TcTV-6)
<i>Crustavirus</i>	<i>Wenzhou crustavirus</i> *	Wēnzhōu crab virus 1 (WzCV-1)
<i>Wastrivirus</i>	<i>Sanxia wastrivirus</i> *	Sānxíá water strider virus 4 (SxWSV-4)

* Asterisks denote type species. [†] Please note that viruses are real objects that are assigned to concepts that are called taxa. Species, genera, families, and orders are taxa. The taxonomic changes listed here pertain to taxon name changes and do not affect virus names. Taxon names are always italicized and always begin with a capital letter. Virus names, on the other hand, are not italicized and are not capitalized, except if the name or a name component is a proper noun. For educational purposes, this column lists the virus names with their correct (lack of) capitalization.

¹ Includes: Newcastle disease virus (NDV) and pigeon paramyxovirus 1 (PPMV-1); ² synonym: anaconda paramyxovirus; ³ synonym: GH-M74a virus; ⁴ synonym: murine parainfluenza virus 1; ⁵ synonym: bat paramyxovirus; ⁶ synonym: simian virus 5; ⁷ synonym: porcine rubulavirus; ⁸ synonyms: avian pneumovirus, turkey rhinotracheitis virus; ⁹ synonyms: citrus leprosis virus nuclear type, citrus necrotic spot virus; ¹⁰ synonym: Tzipori virus; ¹¹ synonym: wild boar rhabdovirus 1 (WBRV1); ¹² synonyms: Egtved virus, Paralichthys olivaceus rhabdovirus; ¹³ synonym: tobacco stunt virus

Acknowledgements We thank Laura Bollinger (NIH/NIAID Integrated Research Facility at Fort Detrick, Frederick, MD, USA) for critically editing the manuscript.

Members of the International Committee on Taxonomy of Viruses (ICTV) *Bornaviridae* Study Group who coauthored this manuscript: Ralf Dürrwald, Masayuki Horie, Jens H. Kuhn, Norbert Nowotny, Susan L. Payne, Dennis Rubbenstroth, Thomas Briese, Martin Schuemme, Keizo Tomonaga.

Members of the ICTV *Filoviridae* Study Group who coauthored this manuscript: Gaya K. Amarasinghe, Christopher F. Basler, Alexander Bukreyev, Kartik Chandran, Olga Dolnik, John M. Dye, Hideki Ebihara, Pierre Formenty, Roger Hewson, Gary P. Kobinger, Jens H. Kuhn, Eric M. Leroy, Elke Mühlberger, Sergey V. Netesov, Jean L. Patterson, Janusz T. Paweska, Sophie J. Smith, Ayato Takada, Jonathan S. Towner, Viktor E. Volchkov, Victoria Wahl-Jensen.

Members of the ICTV *Mononegavirales* Study Group who coauthored this manuscript: Ralf G. Dietzgen, Jens H. Kuhn, Gael Kurath, Bertus K. Rima, Norbert Nowotny, Andrew J. Easton, Dennis Rubbenstroth, Nikos Vasilakis, Peter J. Walker.

Members of the ICTV *Nyamiviridae* Study Group who coauthored this manuscript: Ralf G. Dietzgen, Elodie Ghedin, Dàohóng Jiāng, Jens H. Kuhn, Nikos Vasilakis, David Wang.

Members of the ICTV *Paramyxoviridae* Study Group who coauthored this manuscript: Peter L. Collins, Andrew J. Easton, Ron A. M. Fouchier, Gael Kurath, Robert A. Lamb, Andrea Maisner, Bertus K. Rima, Paul Rota, Benhur Lee, Ming Li, Lin-Fa Wang.

Members of the ICTV *Rhabdoviridae* Study Group who coauthored this manuscript: Kim R. Blasdell, Charles H. Calisher, Ralf G. Dietzgen, Hideki Kondo, Gael Kurath, David M. Stone, Robert B. Tesh, Noël Tordo, Nikos Vasilakis, Peter J. Walker, Anna E. Whitfield.

Compliance with ethical standards

The views and conclusions contained in this document are those of the authors and should not be interpreted as necessarily representing the official policies, either expressed or implied, of the US Department of the Army, the US Department of Defense, the US Department of Health and Human Services, the Department of Homeland Security (DHS) Science and Technology Directorate (S&T) or of the institutions and companies affiliated with the authors. In no event shall any of these entities have any responsibility or liability for any use, misuse, inability to use, or reliance upon the information contained herein. The US departments do not endorse any products or commercial services mentioned in this publication.

Funding This work was supported in part through Battelle Memorial Institute's prime contract with the US National Institute of Allergy and Infectious Diseases (NIAID) under Contract No. HHSN272200700016I. An employee of Battelle Memorial Institute is: A. Bochnowski. A subcontractor to Battelle Memorial Institute who performed this work is: J.H.K., an employee of Tunnell Government Services, Inc. This work was also funded in part under Contract No. HSHQDC-15-C-00064 awarded by DHS S&T for the management and operation of the National Biodefense Analysis and Countermeasures Center (NBACC), a Federally Funded Research and Development Center (V.W.-J.); and National Institutes of Health (NIH) contract HHSN272201000040I/HHSN27200004/D04 and Grant R24AI120942 (N.V., R.B.T.). Y.B. was supported by the Intramural Research Program of the NIH, National Library of Medicine.

Conflict of interest The authors have no conflicts of interest.

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

References

- Adams MJ, Lefkowitz EJ, King AM, Carstens EB (2014) Ratification vote on taxonomic proposals to the International Committee on Taxonomy of Viruses (2014). *Arch Virol* 159:2831–2841
- Afonso CL, Amarasinghe GK, Bányai K, Bào Y, Basler CF, Bavari S, Bejerman N, Blasdell KR, Briand F-X, Briese T, Bukreyev A, Calisher CH, Chandran K, Chéng J, Clawson AN, Collins PL, Dietzgen RG, Dolnik O, Domier LL, Dürrrwald R, Dye JM, Easton AJ, Ebihara H, Farkas SL, Freitas-Astúa J, Formenty P, Fouchier RA, Fù Y, Ghedin E, Goodin MM, Hewson R, Horie M, Hyndman TH, Jiāng D, Kitajima EW, Kobinger GP, Kondo H, Kurath G, Lamb RA, Lenardon S, Leroy EM, Li C-X, Lin X-D, Liú L, Longdon B, Marton S, Maisner A, Mühlberger E, Netesov SV, Nowotny N, Patterson JL, Payne SL, Paweska JT, Randall RE, Rima BK, Rota P, Rubbenstroth D, Schwemmle M, Shi M, Smither SJ, Stenglein MD, Stone DM, Takada A, Terregino C, Tesh RB, Tian J-H, Tomonaga K, Tordo N, Towner JS, Vasilakis N, Verbeek M, Volchkov VE, Wahl-Jensen V, Walsh JA, Walker PJ, Wang D, Wang L-F, Wetzel T, Whitfield AE, Xiè JT, Yuen K-Y, Zhang Y-Z, Kuhn JH (2016) Taxonomy of the order *Mononegavirales*: update 2016. *Arch Virol* 161:2351–2360
- Albariño CG, Foltzer M, Towner JS, Rowe LA, Campbell S, Jaramillo CM, Bird BH, Reeder DM, Vodzak ME, Rota P, Metcalfe MG, Spiropoulou CF, Knust B, Vincent JP, Frace MA, Nichol ST, Rollin PE, Ströher U (2014) Novel paramyxovirus associated with severe acute febrile disease, South Sudan and Uganda, 2012. *Emerg Infect Dis* 20:211–216
- Allison AB, Mead DG, Palacios GF, Tesh RB, Holmes EC (2014) Gene duplication and phylogeography of North American members of the Hart Park serogroup of avian rhabdoviruses. *Virology* 448:284–292
- Baker KS, Todd S, Marsh GA, Crameri G, Barr J, Kamins AO, Peel AJ, Yu M, Hayman DT, Nadim B, Mtové G, Amos B, Reyburn H, Nyarko E, Suu-Ire R, Murcia PR, Cunningham AA, Wood JL, Wang L-F (2013) Novel, potentially zoonotic paramyxoviruses from the African straw-colored fruit bat *Eidolon helvum*. *J Virol* 87:1348–1358
- Binger T, Annan A, Drexler JF, Müller MA, Kallies R, Adankwah E, Wollny R, Kopp A, Heidemann H, Dei D, Agya-Yao FC, Junglen S, Feldt T, Kurth A, Oppong S, Adu-Sarkodie Y, Drosten C (2015) A novel rhabdovirus isolated from the straw-colored fruit bat *Eidolon helvum*, with signs of antibodies in swine and humans. *J Virol* 89:4588–4597
- Bishop DHL, Pringle CR (1995) Order *Mononegavirales*. In: Murphy FA, Fauquet CM, Bishop DHL, Ghabrial SA, Jarvis AW, Martelli GP, Mayo MA, Summers MD (eds) Virus Taxonomy—Sixth Report of the International Committee on Taxonomy of Viruses/Archives of Virology Supplement 10. Springer-Verlag, Vienna, Austria, pp 265–267
- Blasdell KR, Voysey R, Bulach DM, Trinidad L, Tesh RB, Boyle DB, Walker PJ (2012) Malakal virus from Africa and Kimberley virus from Australia are geographic variants of a widely distributed ephemerovirus. *Virology* 433:236–244
- Blasdell KR, Widen SG, Diviney SM, Firth C, Wood TG, Guzman H, Holmes EC, Tesh RB, Vasilakis N, Walker PJ (2014) Koolpinyah and Yata viruses: two newly recognised ephemeroviruses from tropical regions of Australia and Africa. *Vet Microbiol* 174:547–553
- Blasdell KR, Guzman H, Widen SG, Firth C, Wood TG, Holmes EC, Tesh RB, Vasilakis N, Walker PJ (2015) *Ledantevirus*: a proposed new genus in the *Rhabdoviridae* has a strong ecological association with bats. *Am J Trop Med Hyg* 92:405–410
- Burroughs AL, Tachedjian M, Crameri G, Durr PA, Marsh GA, Wang L-F (2015) Complete genome sequence of Teviot paramyxovirus, a novel rubulavirus isolated from fruit bats in Australia. *Genome Announc* 3:e00115–e00177
- Chua KB, Wang L-F, Lam SK, Eaton BT (2002) Full length genome sequence of Tioman virus, a novel paramyxovirus in the genus *Rubulavirus* isolated from fruit bats in Malaysia. *Arch Virol* 147:1323–1348
- Contreras MA, Eastwood G, Guzman H, Popov V, Savit C, Uribe S, Kramer LD, Wood TG, Widen SG, Fish D, Tesh RB, Vasilakis N, Walker PJ (2017) *Almendravirus*: a proposed new genus of rhabdoviruses isolated from mosquitoes in tropical regions of the Americas. *Am J Trop Med Hyg* 96:100–109
- de Almeida Medeiros DB, Diniz Júnior JAP, Cardoso JF, Silva SP, da Silva DEA, de Oliveira LF, é Vasconcelos JM, Chiang JO, Dias AA, Nunes MR, de Viana Júnior SGJL, Vasconcelos PFC (2014) Nearly complete genome sequence of Curionopolis virus, a *Culicoides*-related rhabdovirus isolated in the Brazilian Amazon region. *Genome Announc* 2:e01114–e01158
- Diniz JAP, Nunes MRT, Travassos da Rosa APA, Cruz ACR, de Souza W, Medeiros DBA, Chiang JO, Vasconcelos PFC (2006) Characterization of two new rhabdoviruses isolated from midges (*Culicoides* spp) in the Brazilian Amazon: proposed members of a new genus, Bracorhabdovirus. *Arch Virol* 151:2519–2527
- Drexler JF, Corman VM, Müller MA, Maganga GD, Vallo P, Binger T, Gloza-Rausch F, Cottontail VM, Rasche A, Yordanov S, Seebens A, Knornshild M, Oppong S, Adu Sarkodie Y, Pongombo C, Lukashev AN, Schmidt-Chanasit J, Stöcker A, Carneiro AJ, Erbar S, Maisner A, Fronhoffs F, Buettner R, Kalko EKV, Kruppa T, Franke CR, Kallies R, Yandoko ERN, Herrler G, Reusken C, Hassanin A, Krüger DH, Matthee S, Ulrich RG,

- Leroy EM, Drosten C (2012) Bats host major mammalian paramyxoviruses. *Nat Commun* 3:796
17. Easton AJ, Pringle CR (2011) Order *Mononegavirales*. In: King AMQ, Adams MJ, Carstens EB, Lefkowitz EJ (eds) Virus Taxonomy—Ninth Report of the International Committee on Taxonomy of Viruses. Elsevier/Academic Press, London, pp 653–657
 18. Ghedin E, Rogers MB, Widen SG, Guzman H, Travassos da Rosa AP, Wood TG, Fitch A, Popov V, Holmes EC, Walker PJ, Vasilakis N, Tesh RB (2013) Kolente virus, a rhabdovirus species isolated from ticks and bats in the Republic of Guinea. *J Gen Virol* 94:2609–2615
 19. Goraichuk I, Sharma P, Stegniy B, Muzyka D, Pantin-Jackwood MJ, Gerilovich A, Solodankin O, Bolotin V, Miller PJ, Dimitrov KM, Afonso CL (2016) Complete genome sequence of an avian paramyxovirus representative of putative new serotype 13. *Genome Announc* 4:e00716–e00729
 20. Grard G, Fair JN, Lee D, Slikas E, Steffen I, Muyembe J-J, Sittler T, Veeraraghavan N, Ruby JG, Wang C, Makwana M, Mulembakani P, Tesh RB, Mazet J, Rimoin AW, Taylor T, Schneider BS, Simmons G, Delwart E, Wolfe ND, Chiu CY, Leroy EM (2012) A novel rhabdovirus associated with acute hemorrhagic fever in Central Africa. *PLoS Pathog* 8:e1002924
 21. Gubala AJ, Proll DF, Barnard RT, Cowled CJ, Crameri SG, Hyatt AD, Boyle DB (2008) Genomic characterisation of Wongabel virus reveals novel genes within the *Rhabdoviridae*. *Virology* 376:13–23
 22. Higgins CM, Bejerman N, Li M, James AP, Dietzgen RG, Pearson MN, Revill PA, Harding RM (2016) Complete genome sequence of Colocasia bobone disease-associated virus, a putative cytorhabdovirus infecting taro. *Arch Virol* 161:745–748
 23. Hoffmann B, Tappe D, Höper D, Herden C, Boldt A, Mawrin C, Niedersträßer O, Müller T, Jenckel M, van der Grinten E, Lutter C, Abendroth B, Teifke JP, Cadar D, Schmidt-Chanasit J, Ulrich RG, Beer M (2015) A variegated squirrel bornavirus associated with fatal human encephalitis. *N Engl J Med* 373:154–162
 24. Karamendin K, Kydyrmanov A, Seidalina A, Asanova S, Sayatov M, Kasymbekov E, Khan E, Daulbayeva K, Harrison SM, Carr IM, Goodman SJ, Zhumatov K (2016) Complete genome sequence of a novel avian paramyxovirus (APMV-13) isolated from a wild bird in Kazakhstan. *Genome Announc* 4:e00116–e00167
 25. Kuhn JH, Bekal S, Cai Y, Clawson AN, Domier LL, Herrel M, Jahrling PB, Kondo H, Lambert KN, Mihindukulasuriya KA, Nowotny N, Radoshitzky SR, Schneider U, Staeheli P, Suzuki N, Tesh RB, Wang D, Wang L-F, Dietzgen RG (2013) *Nyamiviridae*: proposal for a new family in the order *Mononegavirales*. *Arch Virol* 158:2209–2226
 26. Lau SKP, Woo PCY, Wong BHL, Wong AYP, Tsui H-W, Wang M, Lee P, Xu H, Poon RWS, Guo R, Li KSM, Chan K-H, Zheng B-J, Yuen K-Y (2010) Identification and complete genome analysis of three novel paramyxoviruses, Tuhoku virus 1, 2 and 3, in fruit bats from China. *Virology* 404:106–116
 27. Lauck M, Yú SQ, Cai Y, Hensley LE, Chiu CY, O'Connor DH, Kuhn JH (2015) Genome sequence of Bivens Arm virus, a tibrovirus belonging to the species *Tibrogargan virus* (*Mononegavirales: Rhabdoviridae*). *Genome Announc* 3:e00015–e00089
 28. Li C-X, Shi M, Tian J-H, Lin X-D, Kang Y-J, Chen L-J, Qin X-C, Xu J, Holmes EC, Zhang Y-Z (2015) Unprecedented genomic diversity of RNA viruses in arthropods reveals the ancestry of negative-sense RNA viruses. *Elife* 4:e05378
 29. McAllister J, Gauci PJ, Mitchell IR, Boyle DB, Bulach DM, Weir RP, Melville LF, Davis SS, Gubala AJ (2014) Genomic characterisation of Almpiwar virus, Harrison Dam virus and Walkabout Creek virus; three novel rhabdoviruses from northern Australia. *Virology Reports* 3–4:1–17
 30. Ng TF, Driscoll C, Carlos MP, Prioleau A, Schmieder R, Dwivedi B, Wong J, Cha Y, Head S, Breitbart M, Delwart E (2013) Distinct lineage of vesiculovirus from big brown bats, United States. *Emerg Infect Dis* 19:1978–1980
 31. Pfeffer M, Dilcher M, Tesh RB, Hufert FT, Weidmann M (2013) Genetic characterization of Yug Bogdanovac virus. *Virus Genes* 46:201–202
 32. Pringle CR (1991) Order *Mononegavirales*. In: Francki RIB, Fauquet CM, Knudson DL, Brown F (eds) Classification and Nomenclature of Viruses—Fifth Report of the International Committee on Taxonomy of Viruses/Archives of Virology Supplementum 2. Springer-Verlag, Vienna, Austria, pp 239–241
 33. Pringle CR, Alexander DJ, Billeter MA, Collins PL, Kingsbury DW, Lipkind MA, Nagai Y, Orvell C, Rima B, Rott R, ter Meulen V (1991) The order *Mononegavirales*. *Arch Virol* 117:137–140
 34. Pringle CR (1996) Virus taxonomy 1996—a bulletin from the Xth International Congress of Virology in Jerusalem. *Arch Virol* 141:2251–2256
 35. Pringle CR (1997) The order *Mononegavirales*—current status. *Arch Virol* 142:2321–2326
 36. Pringle CR (2000) Order *Mononegavirales*. In: van Regenmortel MHV, Fauquet CM, Bishop DHL, Carstens EB, Estes MK, Lemon SM, Maniloff J, Mayo MA, McGeoch DJ, Pringle CR, Wickner RB (eds) Virus Taxonomy—Seventh Report of the International Committee on Taxonomy of Viruses. Academic Press, San Diego, California, USA, pp 525–530
 37. 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
 38. Sakai K, Hagiwara K, Omatsu T, Hamasaki C, Kuwata R, Shimoda H, Suzuki K, Endoh D, Nagata N, Nagai M, Katayama Y, Oba M, Kurane I, Saijo M, Morikawa S, Mizutani T, Maeda K (2015) Isolation and characterization of a novel rhabdovirus from a wild boar (*Sus scrofa*) in Japan. *Vet Microbiol* 179:197–203
 39. Stremlau MH, Andersen KG, Folarin OA, Grove JN, Odia I, Ehiane PE, Omoniwa O, Omologe O, Jiang P-P, Yozwiak NL, Matranga CB, Yang X, Gire SK, Winnicki S, Tariyal R, Schaffner SF, Okokhere PO, Okogbenin S, Akpede GO, Asogun DA, Agbonlahor DE, Walker PJ, Tesh RB, Levin JZ, Garry RF, Sabeti PC, Happi CT (2015) Discovery of novel rhabdoviruses in the blood of healthy individuals from West Africa. *PLoS Negl Trop Dis* 9:e0003631
 40. 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
 41. Vasilakis N, Widen S, Mayer SV, Seymour R, Wood TG, Popov V, Guzman H, Travassos da Rosa APA, Ghedin E, Holmes EC, Walker PJ, Tesh RB (2013) Niakha virus: a novel member of the family *Rhabdoviridae* isolated from phlebotomine sandflies in Senegal. *Virology* 444:80–89
 42. Vasilakis N, Widen S, Travassos da Rosa APA, Wood TG, Walker PJ, Holmes EC, Tesh RB (2013) Malpais Spring virus is a new species in the genus *Vesiculovirus*. *Virol J* 10:69
 43. Vasilakis N, Castro-Llanos F, Widen SG, Aguilar PV, Guzman H, Guevara C, Fernandez R, Auguste AJ, Wood TG, Popov V, Mundal K, Ghedin E, Kochel TJ, Holmes EC, Walker PJ, Tesh RB (2014) Arboretum and Puerto Almendras viruses: two novel rhabdoviruses isolated from mosquitoes in Peru. *J Gen Virol* 95:787–792

44. Walker PJ, Firth C, Widen SG, Blasdell KR, Guzman H, Wood TG, Paradkar PN, Holmes EC, Tesh RB, Vasilakis N (2015) Evolution of genome size and complexity in the *Rhabdoviridae*. PLoS Pathog 11:e1004664
45. Wang F, Fang Q, Wang B, Yan Z, Hong J, Bao Y, Kuhn JH, Werren JH, Song Q, Ye G (2017) A novel negative-stranded RNA virus mediates sex ratio in its parasitoid host. PLoS Pathog 3:e1006201
46. World Organisation for Animal Health (OIE) (2010) Chapter 10.13. Newcastle disease. Terrestrial Animal Health Code, Paris, France. http://web.oie.int/eng/normes/mcode/en_chapitre_1.10.13.pdf
47. Yamamoto E, Ito H, Tomioka Y, Ito T (2015) Characterization of novel avian paramyxovirus strain APMV/Shimane67 isolated from migratory wild geese in Japan. J Vet Med Sci 77:1079–1085