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ICTV VIRUS TAXONOMY PROFILES

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ICTV Virus Taxonomy Profile: Filoviridae

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Abstract

Members of the family *Filoviridae* produce variously shaped, often filamentous, enveloped virions containing linear non-segmented, negative-sense RNA genomes of 15–19 kb. Several filoviruses (e.g., Ebola virus) are pathogenic for humans and are highly virulent. Several filoviruses infect bats (e.g., Marburg virus), whereas the hosts of most other filoviruses are unknown. This is a summary of the International Committee on Taxonomy of Viruses (ICTV) Report on *Filoviridae*, which is available at www.ictv.global/report/filoviridae.

Table 1. Characteristics of members of the family Filoviridae

Typical member:	Marburg virus (DQ217792), species Marburg marburgvirus, genus Marburgvirus
Virion	Enveloped, variously shaped but predominantly filamentous, typically with a single nucleocapsid
Genome	Approximately 15-19 kb of linear, negative-sense, non-segmented RNA
Replication	Antigenomic RNA is a replication intermediate. The genome and antigenome form ribonucleoprotein complexes, which serve as templates
Translation	From multiple 5'-capped and 3'-polyadenylated mRNAs
Host range	Primates (e.g., ebolaviruses, marburgviruses), bats (e.g., marburgviruses), domestic pigs (e.g., Reston virus) and probably fish (e.g., striaviruses, thamnoviruses) become naturally infected
Taxonomy	Realm <i>Riboviria</i> , phylum <i>Negarnaviricota</i> , subphylum <i>Haploviricotina</i> , class <i>Monjiviricetes</i> , order <i>Mononegavirales</i> ; family includes multiple genera

VIRION

Virions are enveloped and diverse in shape and can appear as branched, toroid, U- or 6-shaped, and long filamentous forms (Table 1, Fig. 1). Virions contain ribonucleoprotein (RNP) complexes composed of genomic RNA and, typically, the structural proteins nucleoprotein (NP), polymerase cofactor (VP35), transcriptional activator (VP30), RNP-associated protein (VP24) and RNA-dependent RNA polymerase (L). The matrix protein (VP40) forms a regular layer beneath the viral envelope. Surface spikes formed by glycoproteins ($GP_{1,2}$) are approximately 7 nm in diameter and

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Abbreviations: L, RNA-dependent RNA polymerase; NP, nucleoprotein; RNP, ribonucleoprotein.

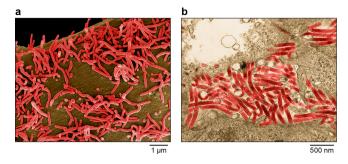


Fig. 1. Electron microscopic images of Marburg virus particles budding from infected Vero E6 cells, (a) scanning EM, (b) transmission EM. Images are colourized for clarity. Courtesy of John G. Bernbaum and Jiro Wada, IRF-Frederick.

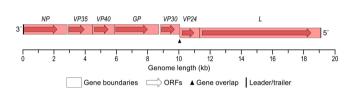


Fig. 2. Schematic representation of the genome organization of Marburg virus. Courtesy of Jiro Wada, IRF-Frederick.

cover the virion surface at approximately 10-nm intervals [1–3]. Some filoviruses do not have discernable glycoproteins and may have different RNP complexes [4].

GENOME

Filovirus genomes are approximately 15–19 kb (Fig. 2) without a 5′-cap or 3′-poly(A). Terminal leader and trailer sequences contain the replication and transcription promoters. Marburg-virus genomes contain seven separate, continuous open reading frames (ORFs) flanked by 3′- and 5′-terminal non-coding regions that contain transcription initiation and termination sites. These ORFs encode the virion structural proteins. Cueva-virus and ebolavirus genomes encode homologues of the marburgvirus structural proteins. However, the marburgvirus GP gene encodes only $GP_{1,2}$, whereas the primary expression product of cuevavirus and ebolavirus GP gene transcription is a soluble glycoprotein. Co-transcriptional editing is used to express $GP_{1,2}$ and additional non-structural proteins. Striaviruses and thamnoviruses encode some, but not all, marburgvirus protein homologues and several proteins of unknown function [4, 5].

REPLICATION

Virus proteins are translated from mRNAs that are synthesized by successive, polar transcription from RNP complexes containing genomic RNA. Replication occurs in the cytoplasm through the synthesis of RNP complexes containing antigenomes that are templates for genomic RNA production. Replication and transcription enzymes include L and VP35. VP30 serves as a transcription enhancer for ebolaviruses and

probably cuevaviruses, but its function in marburgvirus infection is less defined. Virion assembly, including acquisition of the $GP_{1,2}$ -containing lipid envelope, occurs by VP40-mediated budding at the plasma membrane [4, 5].

TAXONOMY

Filoviruses form a family in the haploviricotine order *Mononegavirales*. Within this order, filoviruses are most closely related to members of the families *Paramyxoviridae*, *Pneumoviridae* and *Sunviridae*. The family *Filoviridae* includes multiple genera for viruses that differ in geographic and host range and genomic organization.

RESOURCES

Full ICTV Report on the family *Filoviridae*: www.ictv. global/report/filoviridae.

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Conflicts of interest

The authors declare that there are no conflicts of interest.

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