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**Arenaviruses I - The Epidemiology,
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Arenaviruses are unusually interesting in that they occupy both categories.

Arenaviruses cause several human diseases known primarily as the hemorrhagic fevers occurring in South

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and Latin America (Bolivia: Machupo,
Argentina, Junin virus, and Brazil: Sabia
virus) and in Africa (Lassa fever virus).

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Molecular and Cell Biology of
Arenaviruses. [Michael B A Oldstone] --

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Arenaviruses possess single stranded bi-

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segmented RNA genomes. The large (L) genomic segment (~7,200 nt) encodes the viral RNA-dependant RNA polymerase and a zinc-binding protein. The small (S) genomic segment (~3,500 nt) encodes the nucleocapsid protein (N) and the glycoprotein precursor (GPC) in two non-overlapping reading frames of opposite polarities.

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An arenavirus is a bisegmented
ambisense RNA virus that is a member
of the family Arenaviridae. These viruses

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infect rodents and occasionally humans. A class of novel, highly divergent arenaviruses, properly known as reptarenaviruses, have also been discovered which infect snakes to produce inclusion body disease. At least eight arenaviruses are known to cause human disease.

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Arenavirus - Wikipedia

Phylogeny of New World arenaviruses
based on the complete coding
sequences of the small genomic
segment identified an evolutionary
lineage produced by intrasegmental
recombination. Biochemical and
Biophysical Research Communications.
2002; 296(5):1118-1124.

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Old World/New World Arenaviruses | Viral Hemorrhagic ...

All arenaviruses contain a major nucleocapsid-associated protein of molecular weight 60–68,000 with two glycoproteins in the outer viral envelope. These envelope glycoproteins are not primary gene products but arise by

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proteolytic cleavage of a larger, 75,000
molecular weight glycoprotein precursor
polypeptide (GPC).

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Arenavirus - an overview | ScienceDirect Topics

The Arenaviridae are a family of viruses
whose members are generally
associated with rodent-transmitted

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diseases in humans. Each virus usually is associated with a particular rodent host species in which it is maintained.

Arenavirus infections are relatively common in humans in some areas of the world and can cause severe illnesses.

Arenaviridae | Viral Hemorrhagic Fevers (VHFs) | CDC

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Since the subject of arenaviruses was visited by Current Topics in Microbiology and Immunology 14 years ago, enormous advances have been made in this area. The receptor for several arenaviruses, alpha-dystroglycan, was identified, the replication strategy of these viruses was decoded, and application of a reverse genetics system

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for studying viral gene function and viral
biology is well underway.

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Molecular & Computational biology; ... A
host of disease-causing viruses called
arenaviruses lurk in animal populations
in various parts of the world, sometimes
crossing over into humans. When ...

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Scientists create decoy molecule that neutralizes arenaviruses

Arenaviruses are single-stranded ribonucleic acid (RNA) viruses that cause chronic infections in rodents and zoonotically acquired disease in humans through rodent excreta, especially urine. The...

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Arenaviruses: Practice Essentials, In Background, Pathophysiology

Arenaviruses can cause fatal human haemorrhagic fever (HF) diseases for which vaccines and therapies are extremely limited. Both the New World (NW) and Old World (OW) groups of arenaviruses contain HF-causing

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pathogens. Although these two groups share many similarities, important differences with re ...

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Comparative analysis of disease pathogenesis and molecular ...

Nonetheless, many aspects of the epidemiology, molecular epidemiology and evolution of the virus are still not

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fully understood. The natural course of infection is rather complex and further complicates diagnosis, treatment and the implementation of preventive measures aimed to control the disease.

**Epidemiology, molecular
epidemiology and evolution of ...**
Phylogenetic analyses of Z protein gene

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nucleotide sequences and RNA-
dependent RNA polymerase gene
nucleotide sequences grouped Pirital
virus with Pichindé virus (a South
American arenavirus which, like Pirital
virus, does not appear to be pathogenic
for humans) and placed the Pirital-
Pichindé lineage in a sister relationship
to a lineage represented by Guanarito

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virus and the three other arenaviruses
known to cause hemorrhagic fever in
South America.
Microbiology And Immunology

**Phylogeny of the Venezuelan
arenaviruses.**

LCM virus remains the prototype of the
Arenaviridae and is a common infection
of laboratory mice, rats and hamsters.

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Once thought rare in humans there is now increasing evidence of LCM virus being implicated in renal disease and as a complication in organ transplantation.

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