



XIV Argentine Congress of Virology

Virology for the future:
Science, Technology and Health

27-29 May 2026

UCA Auditories - Puerto Madero, CABA, Argentina



Abstracts Submission Guidelines

viroarg.com/en/abstracts



Abstract Submission Guidelines

Last Update: March 11, 2026

Opening: 9 December 2025

Closing: 13 March 2026

General Regulations for Contributed Papers

All submitted works must be original and must not have been previously accepted for publication or published in any national or international scientific journals. Clinical case series with a minimum of three (3) cases will be accepted. Works previously presented at the XLV ANNUAL SCIENTIFIC MEETING OF THE ARGENTINE SOCIETY OF VIROLOGY (SAV) 2025 may also be submitted.

Abstracts must be submitted exclusively via the online platform, which will be available starting December 9, 2025. The **deadline for receipt is March 13, 2026, at 11:59 PM (23:59 h)**.

Applicants may indicate their preferred presentation format (poster or oral communication); however, the final assignment will be determined by the organizing committee following the review process.

Abstracts must be submitted in Spanish, except for those submitted by non-Spanish-speaking authors, in which case they must be written in English.

During the submission process, authors must select a “Primary Thematic Area” and a “Specific Sub-area” that best fit their work, as described in the sections below.

All accepted abstracts will be published in a digital abstract book with a registered ISBN, which will be available on the congress website in due course.

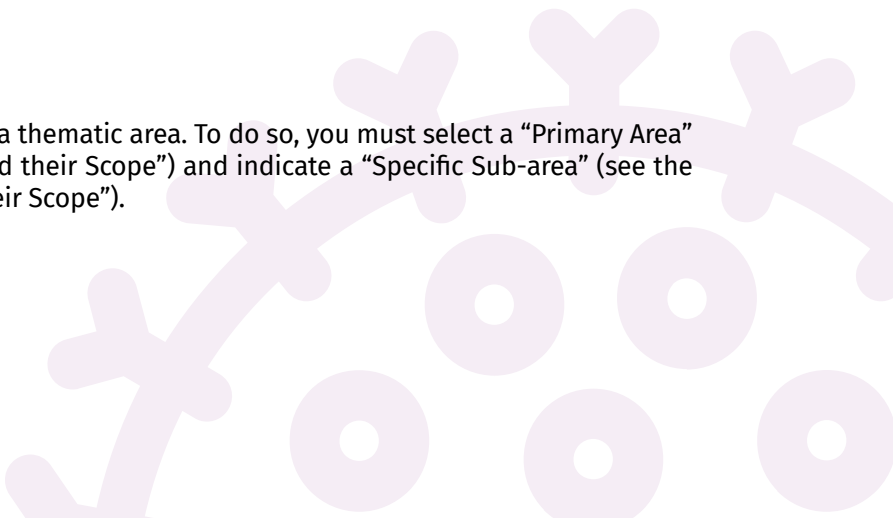
The number of presentations during the event is limited. Therefore, it is recommended not to fragment results from a single experimental line into multiple submissions. Each individual may submit up to one (1) abstract as first author; however, they may participate as co-authors on other works.

Each abstract may have up to eight (8) authors. The individual listed in the first position will serve as the corresponding author and must pay the registration fee once the abstract has been accepted.

The Congress will announce its registration scholarship policy and related regulations in due time.

Thematic Areas: General Overview

Submissions must be assigned to a thematic area. To do so, you must select a “Primary Area” (see the list under “Primary Areas and their Scope”) and indicate a “Specific Sub-area” (see the list under “Specific Sub-areas and their Scope”).





Primary Thematic Areas and their Scope

Below are the eight (8) primary thematic areas to be selected for abstract assignment:

1. Clinical Virology

Studies on viruses that infect humans, including all research and development related to knowledge generation, prevention strategies, diagnostics, and therapeutics. In the case of zoonoses, the study should be included here if the primary focus is on humans.

2. Veterinary Virology

Studies on viruses that infect livestock, companion animals, and wildlife, including all research and development related to knowledge generation, prevention strategies, diagnostics, and therapeutics. In the case of zoonoses, the study should be included here if the primary focus is on animals rather than humans.

3. Environmental and Food Virology

Studies on the presence, distribution, and behavior of viruses in the environment (e.g., water, air, soil, food, among others).

4. Invertebrate Virology

Studies on viruses that infect terrestrial and aquatic invertebrates with productive value (e.g., bees, shellfish) or those considered pests (in agricultural systems or as disease vectors).

5. Prokaryotic Virology

Studies on viruses that infect bacteria or archaea.

6. Plant, Fungal and Protist Virology

Studies on viruses that infect plants, fungi, or protists.

7. Virology and Biotechnology

Developments using virions, their nucleic acids, proteins, or other derived biomolecules for technological purposes in any field of application.

8. Virology and Bioinformatics

Development of computational routines and mathematical models for the analysis of viral nucleotide or protein sequences (or associated host factors), or for the study of epidemiological data regarding viral infections.

Specific Sub-areas and their Scope

The specific thematic areas (listed in alphabetical order) for abstract assignment are detailed below:

A. Antivirals

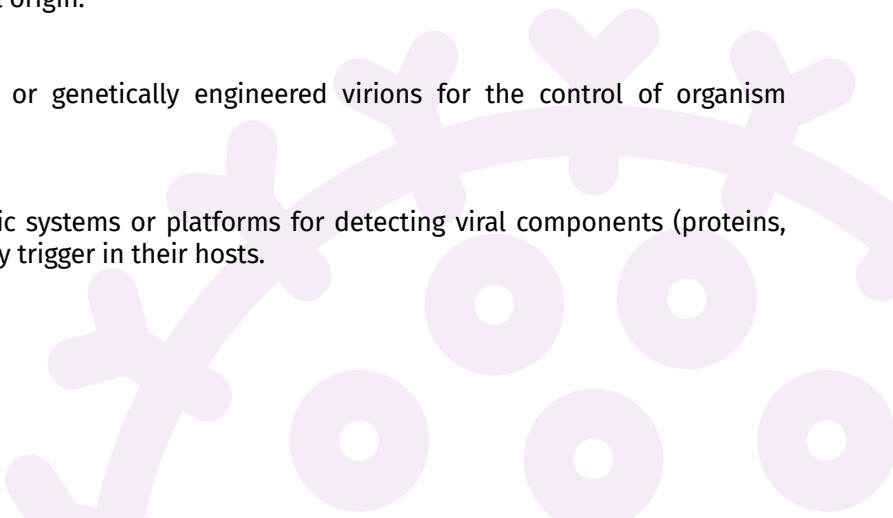
Studies and development of compounds with antiviral activity. This includes both synthetic chemical compounds and those of biological origin.

B. Biological Control

Development and use of natural or genetically engineered virions for the control of organism populations.

C. Diagnostic

Development and use of diagnostic systems or platforms for detecting viral components (proteins, nucleic acids) or the responses they trigger in their hosts.





D.Viral Enzymes

Studies and developments regarding viral enzymes useful in biotechnology and their respective fields of application.

E.Epidemiology

Studies on the distribution, frequency, and patterns associated with viral diseases in host populations. This also includes eco-epidemiological studies that consider the effects of environmental factors.

F.Evolution

Studies on the evolutionary patterns of viral nucleic acids and the proteins they express, as well as studies on host factors linked to viral infections. This includes taxonomic aspects and proposals for modifications in viral classification.

G.Reverse Genetics

Developments and studies based on genetic engineering systems for basic and applied research of viral nucleic acids.

H.Immunity

Studies related to the innate and adaptive immune responses that organisms possess or develop in response to viral infections.

I.Virus-Cell Interactions

Studies related to the interactions between virions and host cells at any stage of the infectious process, excluding immunity-related issues (see specific area "H").

J.Pathogenesis

Studies on the processes involved in the development of viral diseases within host organisms.

K.Replication

Studies related to the components and processes involved in the replication of viral nucleic acids within host cells.

L.Vaccines

Development of new vaccines for the prevention of viral diseases, as well as safety and efficacy studies related to the use of commercially available vaccines.

M.Viral Vectors

Genetic engineering developments based on virions or their nucleic acids for gene delivery to cells and tissues (gene therapy), controlled infection of specific cells (virotherapy), or protein antigen display for vaccine purposes or high-throughput screening.

N.Techniques and methodologies in virology

Experimental technical and methodological developments associated with the handling and study of viruses at any level.

O.Others

Any study or development that cannot be linked to the aforementioned sub-areas.

During the online submission, you may select the Primary Thematic Area and Specific Sub-area, each from its respective drop-down menu.

Each submission must include three (3) keywords. The first one must refer to the main virus under study. The taxonomic name of the viral species (or family/other taxonomic level) must be used. We



recommend visiting the International Committee on Taxonomy of Viruses (ICTV) website (<https://ictv.global>) and reviewing the following documentation:

<https://ictv.global/taxonomy>

<https://www.aam.org.ar/descarga-archivos/virus-nomenclature.pdf>

The other two keywords may refer to specific strains, serotypes, related sub-areas from the provided list, or any other expression directly associated with the focus of the work.

Abstract Format and Upload

It is suggested to draft the abstract in a word processor first and then copy it into the submission form.

Title: Must be precise and not exceed 250 characters (including spaces). It must be written in UPPERCASE; avoid the use of abbreviations. Example: DIFFERENTIAL PROFILE OF EXTRACELLULAR VESICLE-DERIVED MIRNAS IN INDIVIDUALS WITH CHRONIC HEPATITIS C VIRUS INFECTION.

Authorship and Affiliation: Enter full names in the order requested by the form, excluding titles or degrees. Institutional affiliation must be provided for each author following this incremental order: Laboratory or Research Group, Institutional Membership, City, State/Province, Country. Example: Basic Virology Group, Chair of Virology, Institute of Virology, National University of Río Cuarto, Río Cuarto, Córdoba, Argentina. Multiple affiliations must be separated by a semicolon (;). Please note that multiple affiliations will be displayed as shown in the example:

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Abstract Body: Must include Introduction, Objectives, Results, and Conclusions (do not use explicit subheadings for these sections). Bibliographic references, tables, or figures must not be included. If the form allows an attachment (figure/table), it will only be for the evaluation process and will not be included in the abstract book; therefore, they should not be mentioned in the text. The maximum length (excluding title, authors, and affiliations) is 2,500 characters (including spaces). Minimize the use of abbreviations and acronyms; if used, they must be defined upon first mention. Authors are responsible for spelling, syntax, and the correct display of symbols. Proper viral nomenclature must be followed as per the “Keywords” section. Abstracts will be published exactly as submitted. No modifications will be allowed once received by the reviewers.

Evaluation: For “Contributed Papers,” evaluation will be conducted by experts in the selected thematic areas. The decision of acceptance or rejection, along with the presentation format (poster or oral communication), will be communicated to the corresponding author. Accepted works will be required to upload an e-poster to the platform, which will be available to all congress attendees.

Ethical and Environmental Safeguards: By submitting the abstract, authors assume responsibility for complying with all regulations, bioethics standards, and required infrastructure, hygiene, and safety conditions. Works that violate individual identity or gender, involve animal maltreatment, or compromise environmental safety will not be accepted.



Inquiries

For questions regarding scientific or administrative matters related to abstract submission, please contact: cientifica@viroarg.com

Instructions for abstract submission

Open this link

<https://bit.ly/AbstractsSubmission>

and follow the next steps:

- I. The system will prompt you to create an account.**
- II. Use your registered email and password to log in to the system at any time.**
- III. In the “My Abstracts” (Mis Resúmenes) tab, you will find the icon to upload a “New Abstract” on the left-hand side.**
- IV. The following windows will guide you through the submission process. Please complete all required fields.**
- V. Fill in all sections and enter your work in the designated text field.**
- VI. Click “Finish” (Finalizar).**
- VII. The system will display the details of your submitted abstract and send a confirmation copy to your email.**

Accepted Works: Authors of accepted abstracts will be required to upload their e-poster in PDF format to the platform. To do so, log in to the system and go to the “My Presentations” (Mis presentaciones) tab. In the “Actions” (Acciones) column, you will find the icon to attach the .pdf file (maximum size: 9999 MB).

