Genomic Surveillance of AMR across the Human–Animal–Environmental Interface

Abstract Guidelines

We invite abstracts describing completed research, ongoing projects, or project proposals that align with one or more of the following themes:

- **Genomic surveillance** of multidrug-resistant bacteria, fungi, parasites, or viruses
- **Wildlife, food chain, and wastewater** as reservoirs for AMR spread
- **Data standards and ISO accreditation** in genomics
- **Social science and community engagement** in genomic surveillance
- **Policy and ethics in genomics**, including data sharing and repositories

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Abstract Format and Instructions

Title:
Provide a concise and informative title.

Author names and affiliations:
List each author's first name followed by their surname, along with institutional affiliation(s).

Abstract length:
The main text should not exceed **300 words**.

Abstract content:
The abstract should be **clear, concise, and well-structured**, including the following elements as appropriate:

- **Background:** Briefly describe the problem or context being addressed.
- **Methods:** Summarise the experimental approach, data analysis, or framework used.
- **Results:** Present key findings or progress to date.
- **Conclusions:** Highlight the main outcomes, implications, or next steps.

For **project ideas or proposals**, include:

- Background and justification

- Proposed methods
- Preliminary results (if available) or expected outcomes

Please avoid vague statements such as *"A solution to XYZ problem will be presented/discussed."*

Selection Process

Deadline: 8 December 2025, at 23:59 GMT.

The Scientific Committee will review abstracts and select submissions for **oral presentations or posters** based on clarity, relevance, and scientific merit.

References:
A reference list is **not required**.