

Discovery and characterisation of viral biocontrol candidates: viromics contribution to plant protection

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Discovery and characterisation of viral biocontrol candidates: viromics contribution to plant protection

Sarah François

Aymeric Antoine-Lorquin Doriane Mutuel Lisa Claude Marie Frayssinet Mylène Ogliastro



Context

The European legislation to ban many pesticides has led to increased demand for alternatives, including the use of viruses as biocontrol agents, which first requires knowledge of their diversity.

Objectives

Our objectives were to discover insect viral resources, and explore their potential for biocontrol.

Luciana Tavella Luciana Galetto Simona Abba Massimo Turina



Methods

We collected

3 major agricultural pests

and their host-plants

from agricultural ecosystems

located in France and Italy.

We processed

~1000 samples, of pooled or
unpooled individuals,
by viromics* coupled to
an automated pipeline**

for virus identification.

* The term viromics designates the study of viral communities through the without a priori detection and characterization of virus genome sequences.

Our complete virion-associated nucleic acids (VANA) metagenomic protocol is available at:

** The NearVANA pipeline, developed by Aymeric Antoine-Lorquin, is available at:



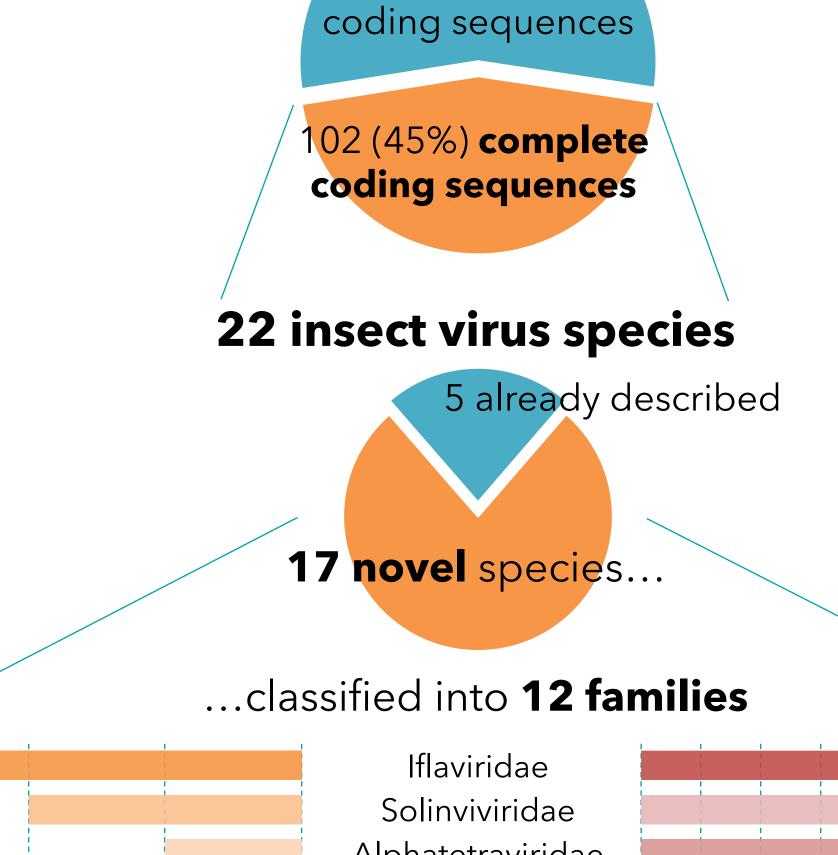


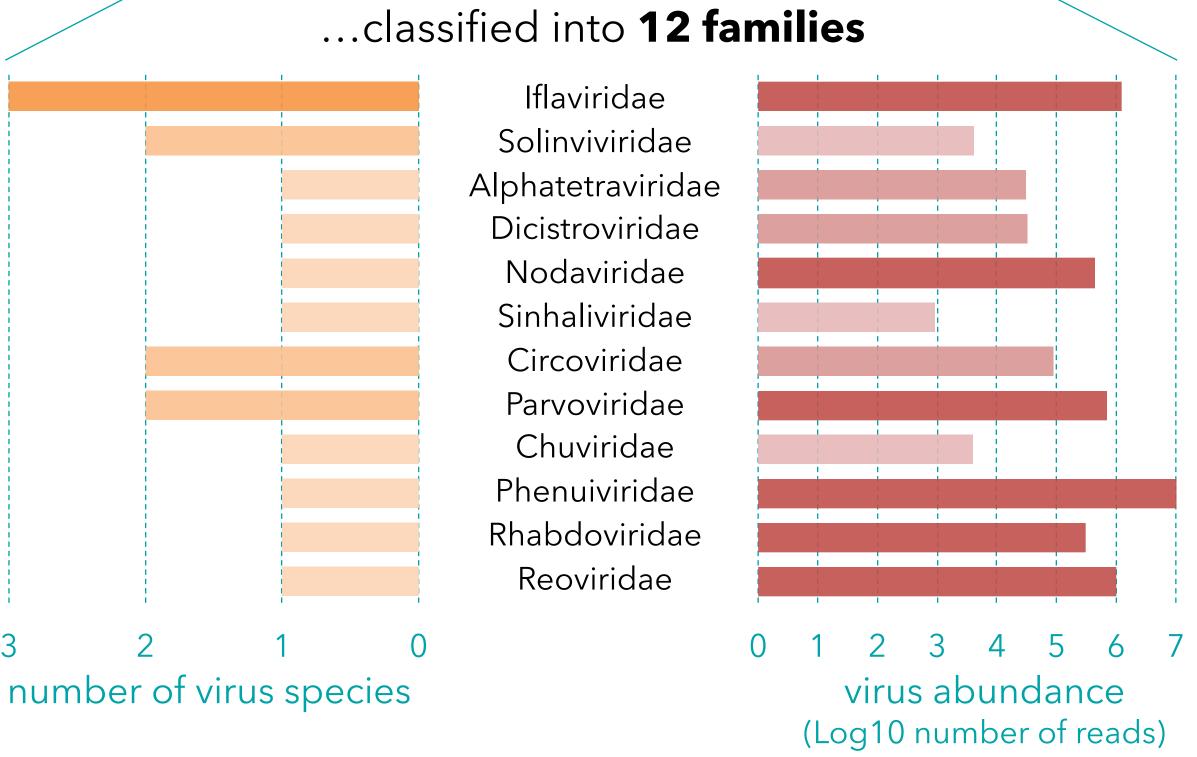


226 virus species

comprising insect viruses, bacteriophages and diet contaminants (plant and fungus viruses)

124 (55%) partial





Results

We revealed a **high diversity** of viruses associated with insect pests:



We detected **54 families of viruses**.



We could classify some of those viruses into 236 virus species*.



We reconstructed the complete coding sequence of 58 novel species, including 17 insect viruses, 26 plant viruses and 12 bacteriophages.



We also showed **disparities**in insect viruses prevalence and
abundance, which are potentially
linked to their **host range**.

* According to the International

Committee on Taxonomy of Viruses (ICTV)

standards

Conclusion & Perspectives

We discovered insect viruses in 3 major agricultural pests.

The characterisation of these viruses (*i.e.* spatio-temporal distribution and phylogenetic analyses) is ongoing.

Their impacts on insect pests will be investigated in future studies.



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