

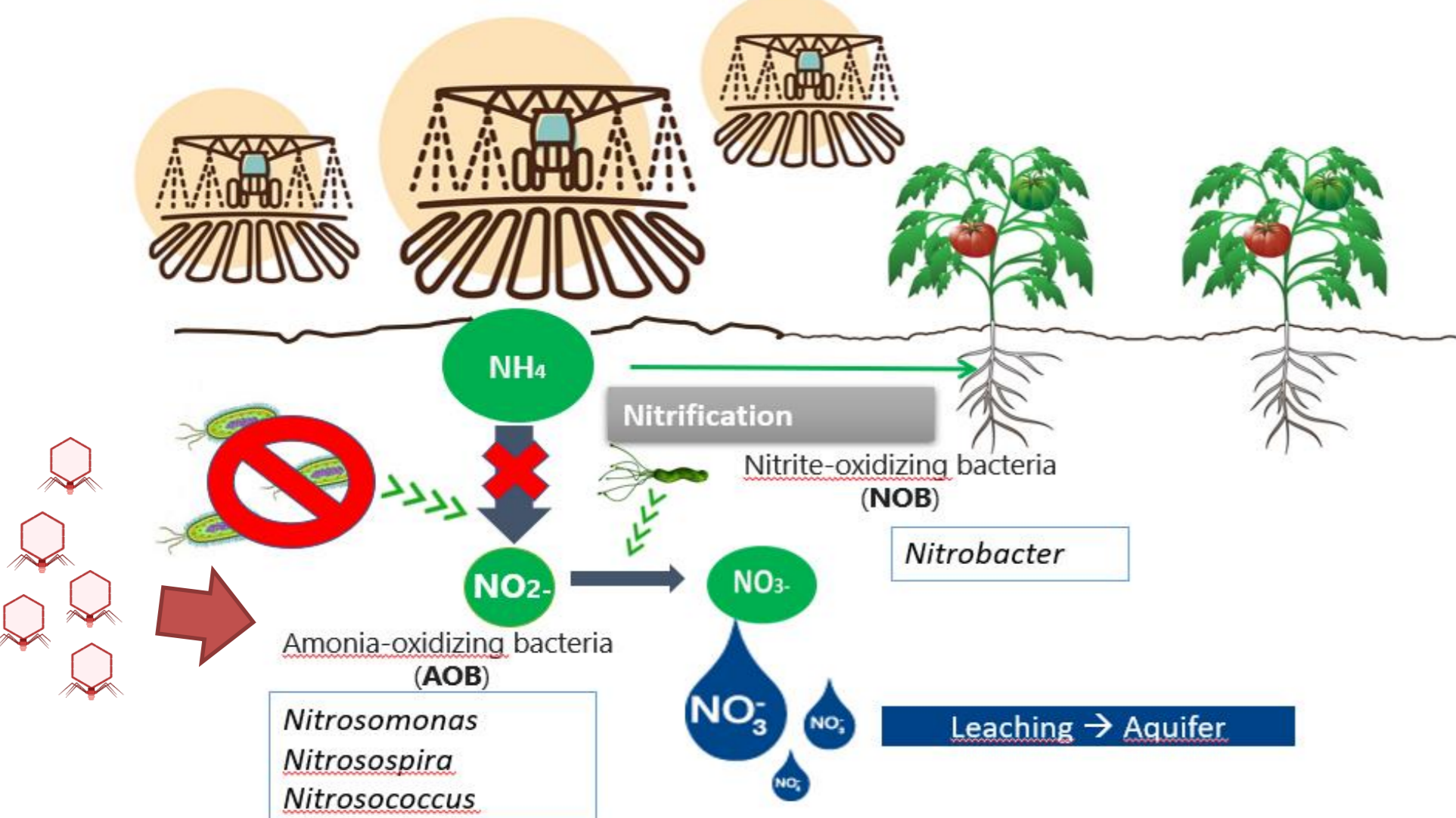
Phage-based biocontrol of nitrification in agricultural soil

Laura Sala-Comorera¹, Pedro Blanco-Picazo¹, Gloria Vique¹, Pablo Quirós², María Dolores Ramos-Barbero¹, Clara Gómez-Gómez¹, Lorena Rodríguez-Rubio¹, Tula Yance-Chávez², Ignasi Salaet², Maite Muniesa¹

1 - Department of Genetics, Microbiology and Statistics. University of Barcelona, Spain
2 - Departamento of I+D+i Fertinagro Biotech S.L., Teruel, Spain

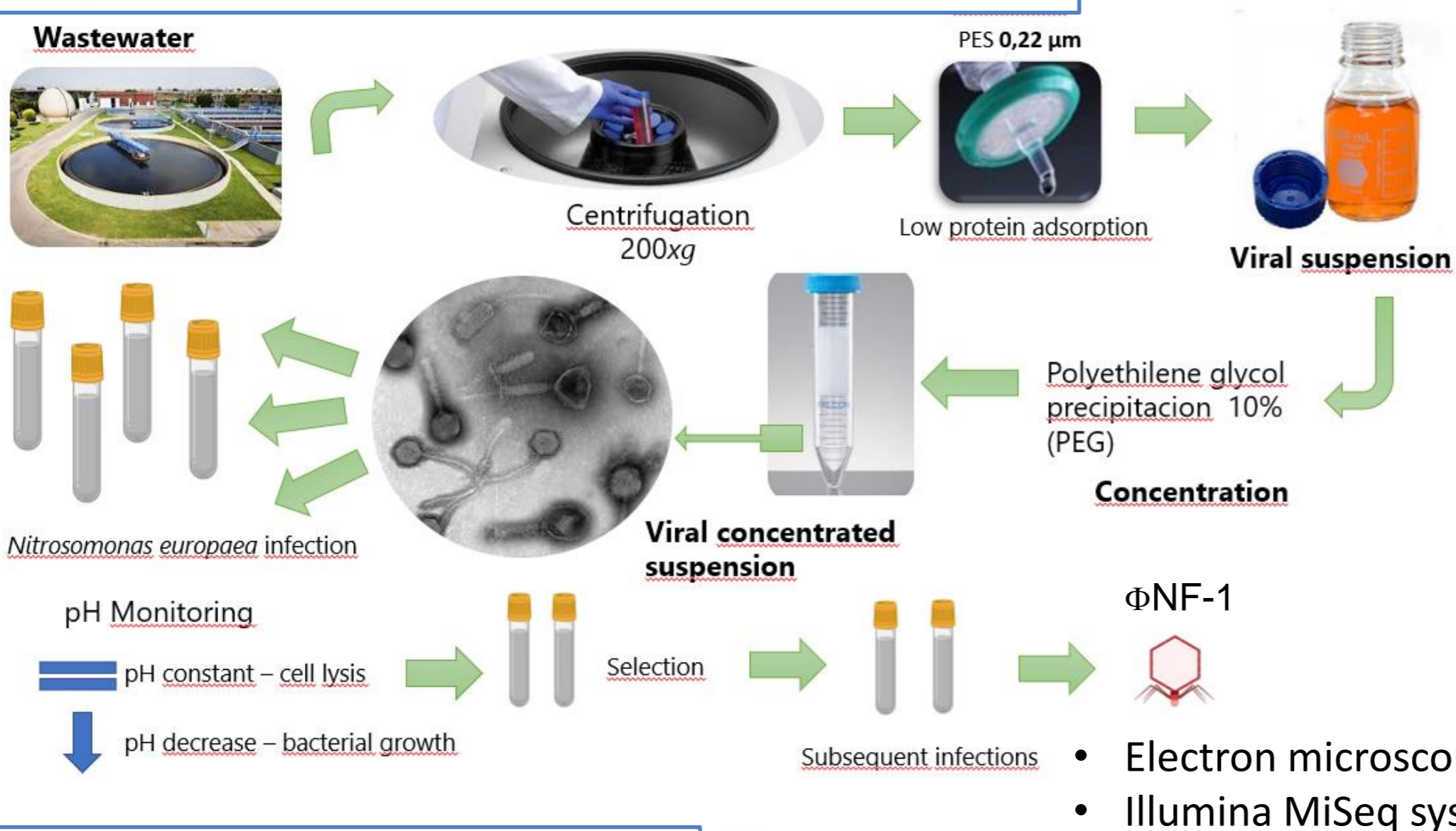
INTRODUCTION & AIM

Nitrogen fertilization boosts crop yields but is inefficient due to the rapid activity of nitrifying bacteria, which leads to the loss of useful nitrogen forms. This requires the use of large amounts of fertilizers, leading to environmental pollution from compounds like NO₃ and N₂O. Although chemical inhibitors can improve fertilization, their impact on the environment and human health remains uncertain. A new strategy is the use of bacteriophages (phages) that specifically target nitrifying bacteria.



METHOD

Phage isolation and characterization



Host spectra and Infectivity

Nitrosomonas and *Nitrospira* sp.

ΦNF-1

In vitro:
AOB medium

In vivo:
soil samples

Availability to infect and inhibit nitrification

- No media acidification
- No ammonium reduction (colorimetric assay)
- No nitrite formation (colorimetric assay)
- Increase in bacteriophage particles (qPCR)

Persistence

pHs 3, 5, 6, 7, 8,

4, 15, 20, 30 °C

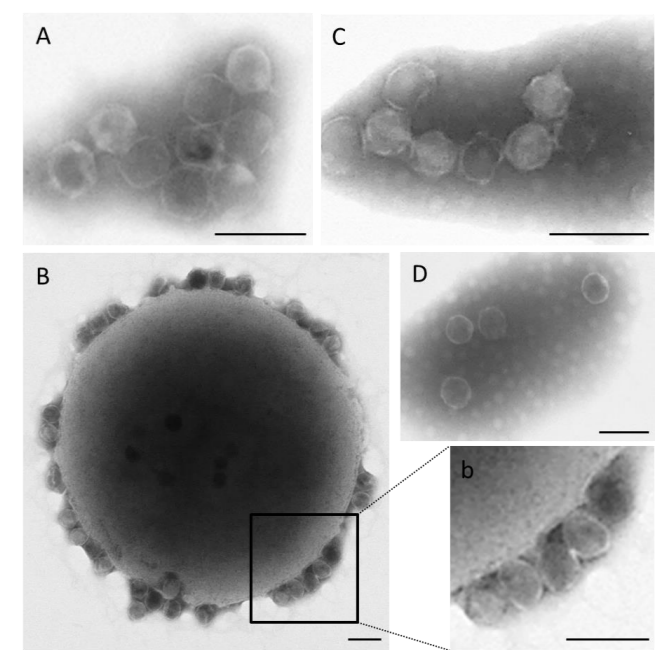
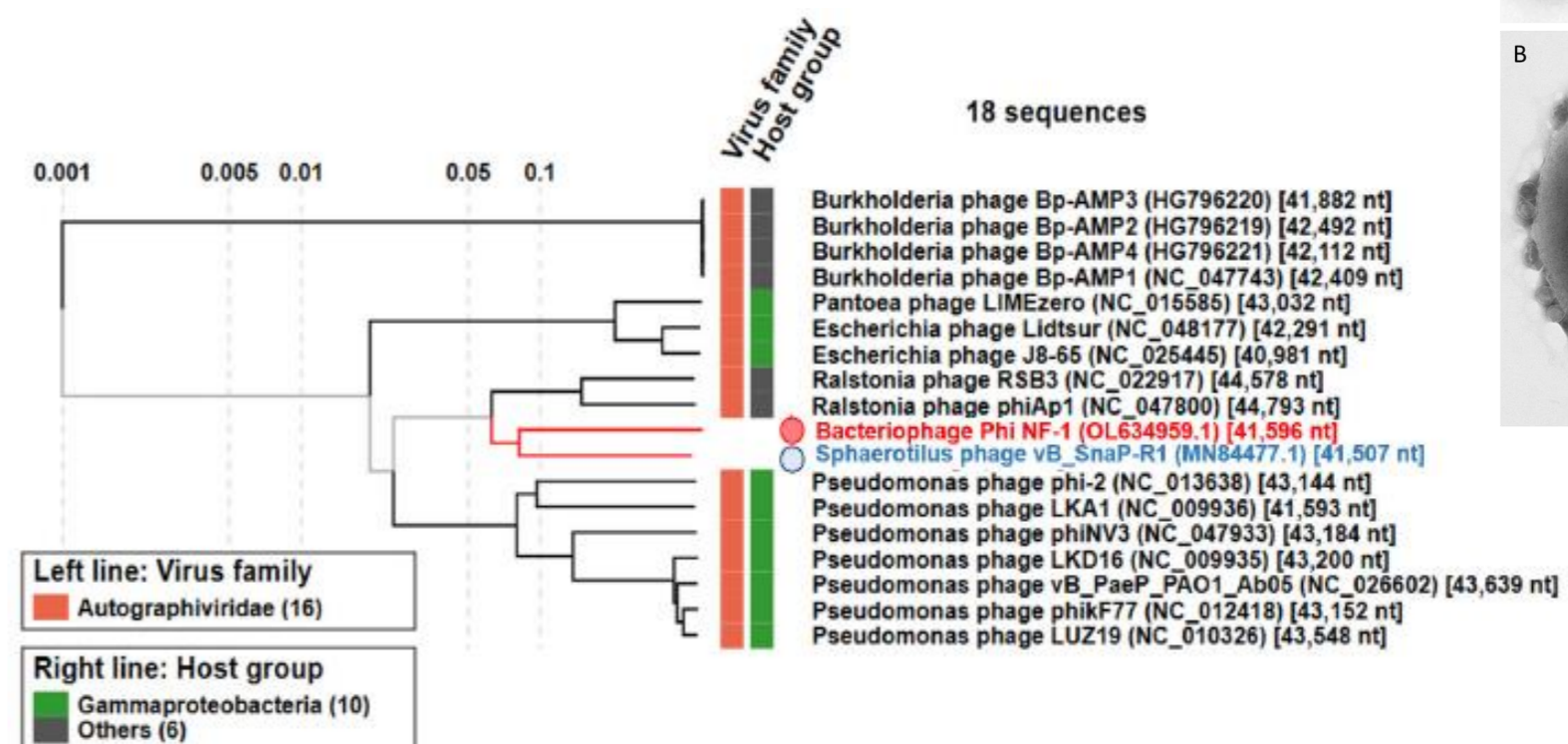
Soil at 20 °C

Concentration of infectious bacteriophage particles over 6 months → Propagation in AOB medium

RESULTS & DISCUSSION

Morphological and Genetic characterization

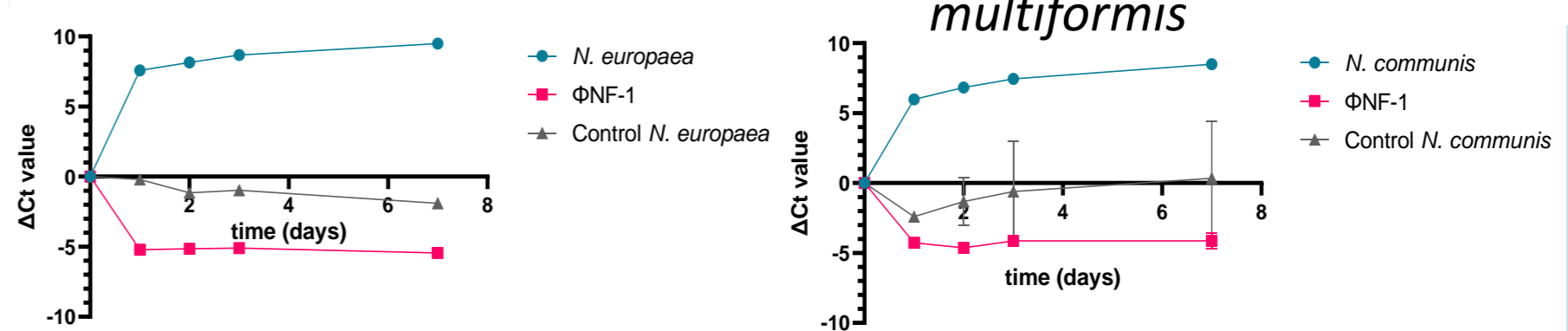
Phage ΦNF1, isolated from wastewater, had an icosahedral capsids of 50 ± 3 nm in diameter and very small tails typical of *Podoviridae* phages.



Phage ΦNF-1 had a genome of 41,596 bp and 45.09% GC content and showed 53 predicted ORFs. The genome analysis of ΦNF-1 suggested it was a virulent phage rather than a temperate one, as it lacks genes associated with lysogen. The most closed sequence was from a *Podoviridae* phage infecting *Sphaerotilus natans*.

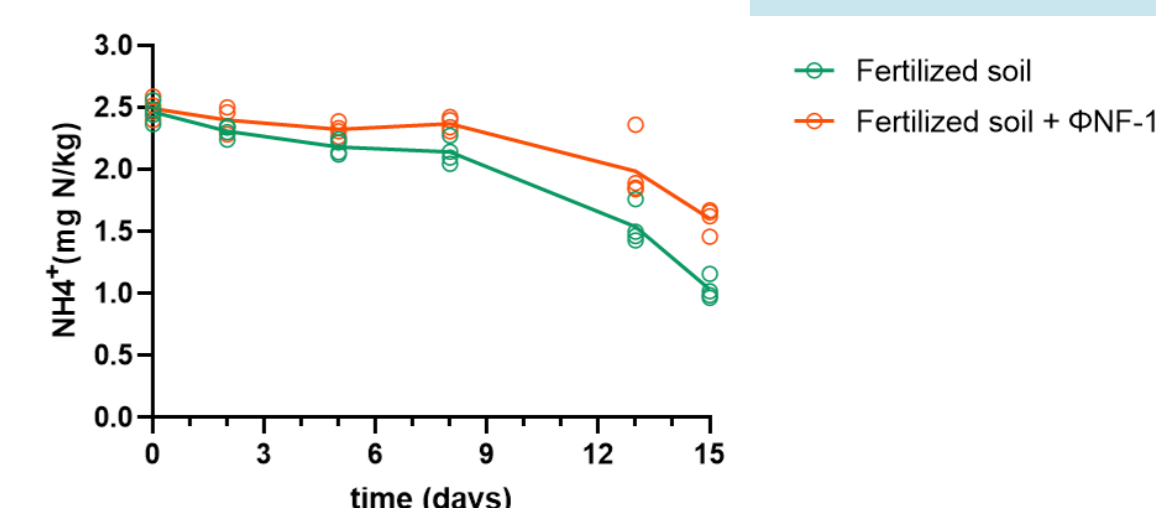
Host spectra and Infectivity

ΦNF-1 was capable of infecting 4 species of *Nitrosomonas* genus and *Nitrospira multififormis*

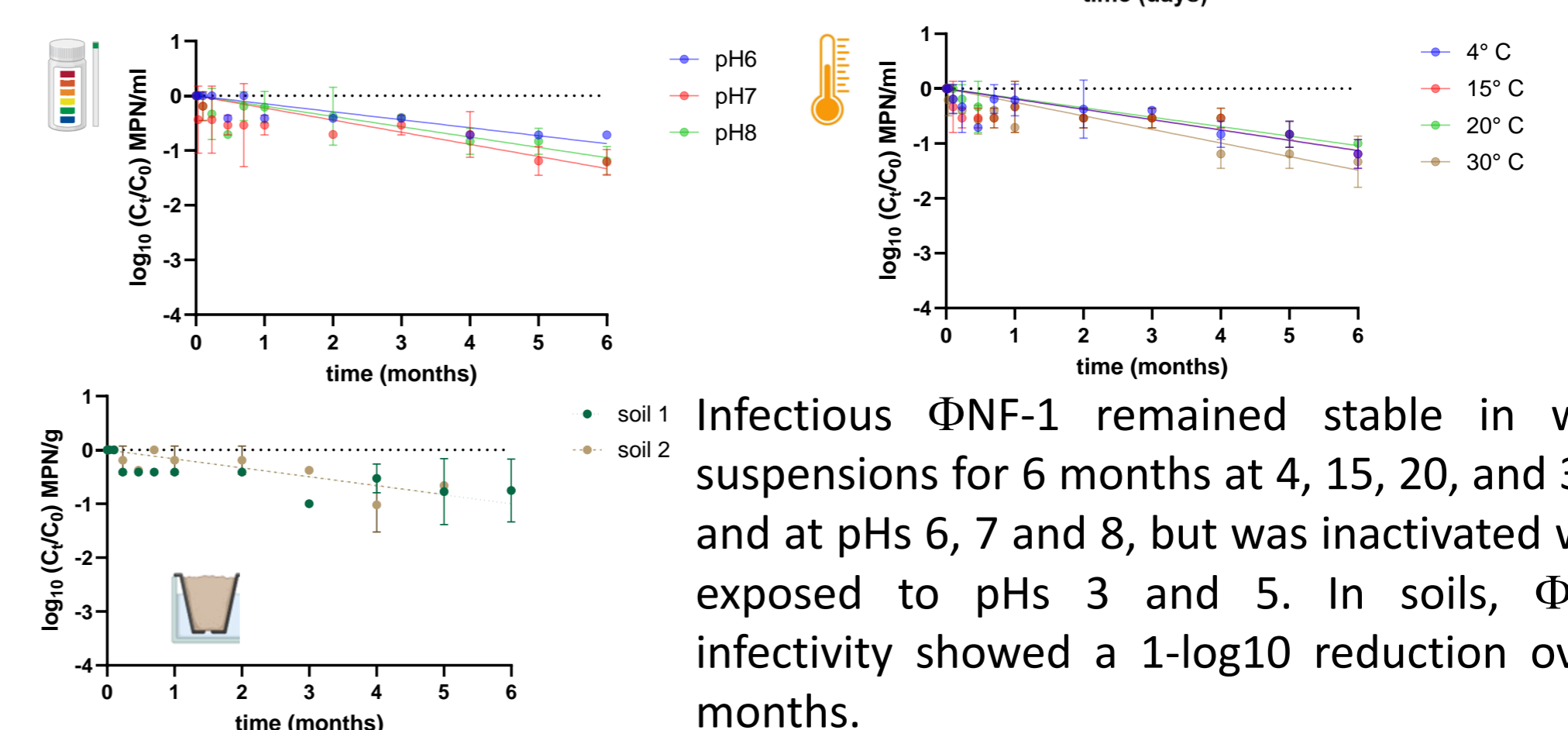


Host spectra-
Nitrosomonas:
N. europaea
N. communis
N. nitrosa
N. oligotropha

ΦNF-1 inhibited bacterial growth, resulting in a reduction of nitrite formation in phage-treated samples in soils and a lower decrease of NH₄⁺



Persistence



CONCLUSION

This study presents a novel approach using bacteriophages to eliminate nitrifying bacteria. Bacteriophages as biocontrol agents offer an effective and environmentally friendly alternative to inhibit nitrification while overcoming the limitations of chemical products.

FUTURE WORK

This application shows potential for improving agricultural fertilization, further research is required to scale up and optimize a product.