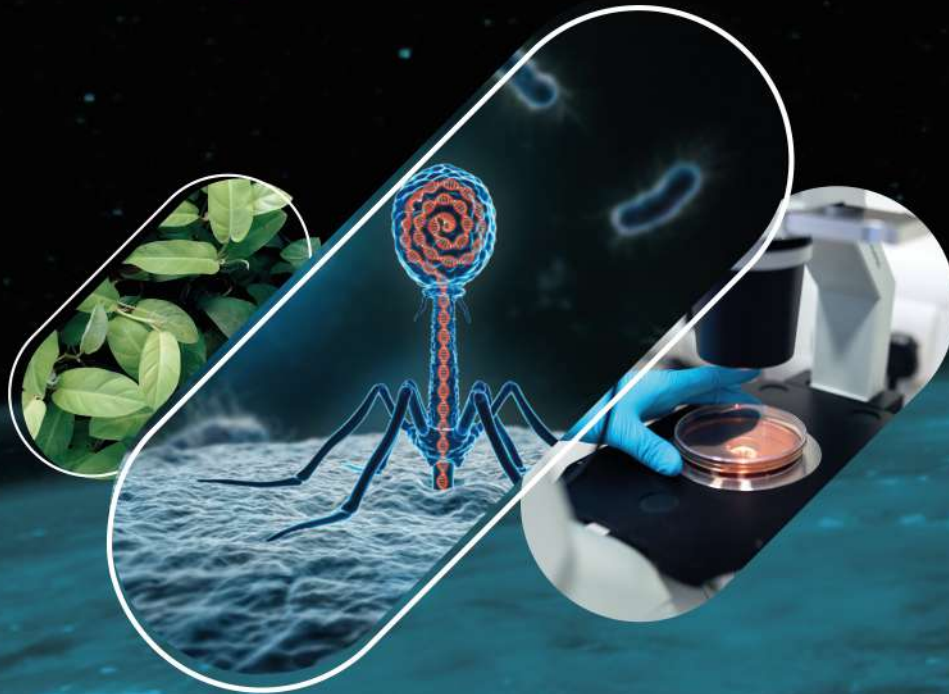


FIRST INTERNATIONAL CONGRESS OF

Bacteriophages and their Applications for the Environment

● *Real world solutions with bacteriophages technology*



 14 - 16 August

ISSN: 2961-2950 [online]

**First International Congress of Bacteriophages
and their Applications for the Environment
(FICBAE)**

© Universidad de Ingeniería y Tecnología, 2023

Jr. Medrano Silva 165, Barranco, Lima, Perú

<https://utec.edu.pe/eventos/congress-of-bacteriophages>

Technical Committee

Alberto Donayre Torres

Mónica Santa-María Fuster

Julio Valdivia Silva

Scientific Committee

Martin Gutierrez Pescarmona

Francisco Salcedo Arancibia

Harry Gustavo Saavedra Espinoza

First Digital Edition: August 2023

Made the Legal Deposit at the National Library
of Peru N° 2023 - 09834

ISSN: 2961-2950 (online)

Design and layout

María De Fátima Salazar Anhuamán

September 2023, Lima - Perú

Contents

Welcome to FICBAE 2023	4
Organizing Committee	5
Topics	6
Program	7
Workshops description	10
Keynote Talks	11
Abstracts	18
Genetic improvement of bacteriophage $\phi XaF13$ for the biological control of bacterial spot in peppers	19
Engineering p8 protein from <i>M13 bacteriophage</i> with high-affinity heavy metal-binding peptides	20
Isolation of lytic Raoultella bacteriophage	21
Isolation and characterization of a lytic bacteriophage isolate from chicken intestine and specific for Multidrug-resistant <i>Salmonella Infantis</i>	22
Morphological and molecular characterization of a lytic bacteriophage isolated from soils from bean crops	23
Isolation of lytic bacteriophages against strains of <i>Klebsiella pneumoniae</i> and <i>Salmonella spp.</i> multidrug-resistant.	24
Scaling of formulated bacteriophages for biological control of bacterial spot (<i>Xanthomonas euvesicatoria</i>) in pepper crops	25
Posters	26
Genetic improvement of bacteriophage $\phi XaF13$ for the biological control of bacterial spot in peppers	27
Engineering p8 protein from <i>M13 bacteriophage</i> with high-affinity heavy metal-binding peptides	28
Isolation of lytic Raoultella bacteriophage	29
Scaling of formulated bacteriophages for biological control of bacterial spot (<i>Xanthomonas euvesicatoria</i>) in pepper crops	30
Workshob lab protocols	31

Welcome to FICBAE 2023

Welcome to The First International Congress of Bacteriophages and their Applications for the Environment (FICBAE). This conference aims to connect with various technology-related universities, agencies, industries, start-ups, and academia to cover topics related to national and international studies of bacteriophages, focusing on efforts for environmental solutions.

Bacteriophages are remarkable viruses that exclusively infect bacteria. These microscopic entities are natural predators of bacteria and are pivotal in shaping microbial ecosystems. The event aims to review the use of bacteriophages for environmental solutions and their impact in Peru and the South America region.

The Congress FICBAE 2023 consists of 4 workshops, 6 keynote talks, 11 specialized talks, and 2 poster sessions. They are organized into 7 topics:

- Phage-based nanomaterials
- Genetic modification of bacteriophages
- Biosecurity
- Bacteriophages in the food industry
- Phage therapy in aquaculture
- Phage therapy in agriculture
- Phage genome analysis

In addition, we have distinguished invited speakers from Peru, Mexico, USA, and Argentina for the keynote seminars. We would like to thank all the researchers, students, and individuals for their contributions. This event is organized by the Bioengineering program and the Bioengineering Research Center, CentroBio, at UTEC University, with the support of PROCENCIA.

Organizing Committee

Technical Committee

Dr. Alberto Donayre Torres
Universidad de Ingeniería y Tecnología ■ ■
adonayre@utec.edu.pe

Dra. Mónica Santa-María Fuster
Universidad de Ingeniería y Tecnología ■ ■
msantamaria@utec.edu.pe

Dr. Julio Valdivia-Silva
Universidad de Ingeniería y Tecnología ■ ■
jvaldivias@utec.edu.pe

Scientific Committee

Dr. Martin Gutierrez Pescarmona
Universidad Diego Portales ■ ■
martin.gutierrez@mail.udp.cl

Dr. Francisco Salcedo Arancibia
Universidad de Guadalajara ■ ■
f.salcedo@cucea.udg.mx

Dr. Harry Gustavo
Universidad de Ingeniería y Tecnología ■ ■
hsaavedra@utec.edu.pe

Topics

Phage-based nanomaterials

Innovative biotechnological materials engineered from bacteriophages, offering versatile applications in fields such as drug delivery and tissue engineering.

Genetic modification of bacteriophages

Involves altering the genetic makeup of these viruses to enhance their effectiveness in targeted bacterial infection control, contributing to advancements in biotechnology and medicine.

Biosecurity

Encompasses strategies and measures aimed at safeguarding biological materials, research, and facilities to prevent misuse or unauthorized access to potentially harmful agents, ensuring the responsible practice of science.

Bacteriophages in the food industry

Being explored as natural antimicrobial agents to combat bacterial contamination, offering a promising solution to improve food safety and quality. Also, bacteriophages are pathogenic agents against lactic acid bacteria from dairy industry and phage monitoring and control is a common practice.

Phage therapy in aquaculture

Involves the use of bacteriophages to control bacterial infections in aquatic organisms, reducing the reliance on antibiotics and promoting sustainable practices in the aquaculture sector.

Phage therapy in agriculture

An emerging approach using bacteriophages to combat plant pathogenic bacteria, offering an environmentally friendly alternative to traditional chemical pesticides.

Phage genome analysis

The process of sequencing and studying the genetic information of bacteriophages, providing insights into their diversity, evolution, and potential applications in various fields of science and biotechnology.

Program

DAY 1

8:45 - 9:15	Words of welcome Dr. Alberto Donayre Torres UTEC - Peru
9:15 - 10:00	Keynote Talk: “Biocontrol of plant diseases through plant phage therapy” Dr. Gabriel Rincón Enríquez Centro de Investigación CIATEJ - Mexico
10:00 - 10:30	Phage infections in the dairy industry Dra. Mariángeles Brigglier Marcó Universidad Nacional del Litoral - Argentina
10:30 - 11:00	Coffee Break
11:00 - 11:30	Environmental phages in foods Dra. Maryoris E. Soto Lopez Universidad de Córdoba - Colombia
11:30 - 12:00	Use of bacteriophages to combat bacterial canker on kiwis Dr. Roberto Bastías Romo Pontificia Universidad Católica de Valparaíso - Chile
12:00 - 12:45	Discussion Panel: Bacteriophages in the environment and proposed solutions - What are the challenges for food and agriculture?
12:45 - 14:00	Lunch
14:00 - 14:15	Words of welcome Javier Bustamante UTEC - Peru
14:15 - 14:30	Venue Dr. Julio Valdivia UTEC - Peru
14:30 - 14:45	Congress Overview (Objectives & MISTI UTEC Program) Dr. Alberto Donayre
14:45 - 15:30	Keynote Talk: “Bacteriophages for real world applications and environmental solutions” Dr. Angela Belcher MIT - USA
15:30 - 16:00	Bacteriophages and heavy metals at UTEC Dr. Alberto Donayre UTEC - Peru
16:00 - 16:30	Coffee Break
16:30 - 17:15	Keynote Talk: “Phages and Biomaterials for bioremediation” Dr. Nimrod Heldman MIT - USA
17:15 - 17:45	Scale-up systems for bacteriophage production Msc. Jay Sacane USA
17:45 - 18:15	Discussion Panel: Bacteriophages and bioremediation with biomaterials
18:15 - 18:30	Poster hanging
18:30 - 19:45	Poster Session

DAY 2

9:30 - 11:00	1st Workshop Using of Molecular Cloning Techniques for phage genome (vector M13KE) editing Dr. Nimrod Heldman MIT - USA
11:00 - 11:30	Coffee Break
11:30 - 14:00	2nd Workshop Amplification of recombinant phages at high scale. Purification using centrifugation and chemical precipitation Dr. Nimrod Heldman MIT - USA
14:00 - 15:30	Lunch
15:30 - 16:10	Keynote Talk: “Bacteriophages in the environment experiences in Perú” Dr. Miguel Talledo UNMSM - Peru
16:10 - 16:40	Detection, isolation and characterization of bacteriophages against pathogens of farmed fish. Dra. Violeta Flores IMARPE - Peru
16:40 - 17:10	Phagotherapy in Aquaculture and experiences at the Instituto del Mar del Perú. Dra. Carla Ivonne Fernandez Espinel IMARPE - Peru
17:10 - 17:50	Keynote Talk: “The Comeback of Phage – Episode II” Dr. Pohl Milón UPC - Peru
17:50 - 18:20	Coffee Break
18:20 - 19:00	Discussion Panel: Experiences with bacteriophages in Peru and regulations for their use
19:00 - 19:45	Poster Session

DAY 3

9:00 - 10:30	3rd Workshop Transformation of phage genome into XL1-Blue cells using chemical transformation Msc. Jay Sacane USA
10:30 - 11:00	Coffee Break
11:00 - 12:30	4th Workshop Titering of M13 bacteriophages for selection of cells expressing phages Quantification of phage concentration using a spectrophotometer Msc. Jay Sacane USA
12:30 - 14:00	Lunch
14:00 - 14:45	Keynote Talk: “Bacteriophages and endolysins active against Staphylococcus aureus: from the counter to the patient” Dr. Hector Morbidoni Argentina
14:45 - 15:20	Genomic characterization and bioinformatics analysis of bacteriophages Dr. Cristian Suarez CONICET - Argentina
15:20 - 16:00	Regulations associated with the use of bacteriophages in Peru: Biosafety and access to genetic resources. Dr. David Eduardo Castro MINAM - Peru
16:00 - 16:30	Coffee Break
16:30 - 17:00	Discussion Panel: Genetic diversity of bacteriophages and genetic manipulations
17:00 - 17:30	Poster Award
18:00 - 19:00	Event Closing

Workshops description

Workshop 1: Using of Molecular Cloning Techniques for phage genome (vector M14KE) editing

Workshop 1 offers an exciting exploration of Molecular Cloning Techniques, focusing on phage genome editing using the M14KE vector. This workshop covers the innovative Gibson assembly technique, enabling seamless DNA fragment assembly with an efficient enzyme mix. Participants will gain hands-on experience with tools to unlock the potential of molecular cloning in research and biotechnology.

Workshop 2: Amplification of recombinant phages at high scale. Purification using centrifugation and chemical precipitation

Workshop 2 immerses participants in high-scale recombinant phage amplification and purification techniques using centrifugation and chemical precipitation. This workshop encompasses key experiences for bacterial selection and gene expression studies, providing practical skills vital for molecular biology and biotechnology applications.

Workshop 3: Transformation of phage genome into XL1-Blue cells using chemical transformation

The workshop begins with an experience centered on phage titering, a technique used to assess bacteriophage concentration. This involves serial dilutions of a genetically transformed bacterial culture and plating on agar plates to visualize phage plaque formation. Participants will gain practical insights into the phage titering process and the subsequent steps for phage plating using Dilution tube 8 ($1 : 10^8$), preparing them for hands-on molecular biology applications.

Workshop 4: Titering of M13 bacteriophages for selection of cells expressing phages Quantification of phage concentration using a spectrophotometer

This workshop covers the process of amplifying and purifying bacteriophages, specifically M13KE. The key steps include creating overnight cultures, amplifying the phages, and conducting purification rounds. The final stage involves quantifying phage concentration using a NanoDrop. This workshop provides essential techniques for producing high-quality phage samples.



Keynote Talks

Biocontrol of plant diseases through plant phage therapy

Dr. Gabriel Rincón Enríquez | Centro de Investigación CIATEJ - Mexico

Video

Plant phage therapy focuses on controlling phytopathogenic microorganisms that can lead to 100% crop losses. Alternatives to chemical antibiotics and antifungals involve biological control, such as the use of bacteriophages, viruses that target bacteria, known for their specificity, abundance, and lack of contamination. Laboratory studies evaluated bacteriophages like Φ XaF18, which demonstrated resistance to UV light in specific formulations (F2 and F4). In greenhouse trials, these phages reduced diseases by up to 95%. Additionally, research into Φ XaF13 aimed to identify genes conferring UV resistance and enhance virulence. Both phages were combined to create “Fagolytic”, a product competitive with commercial antibiotics and antifungals. Production scaled from 0.5L to 100L, with pricing adaptable to the agricultural market.



Figure 1. Fagolytic

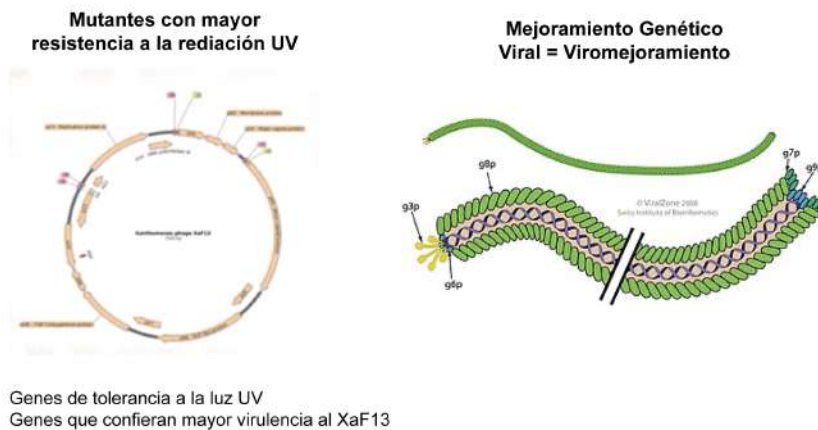


Figure 2. Viral breeding

Bacteriophages for real world applications and environmental solutions

Dr. Angela Belcher | Massachusetts Institute of Technology - USA

Video

Genetic Control of Materials explores how organisms utilize their genetic information to create protective shields and hard materials. This research delves into the potential of leveraging genetic data for the development of batteries and solar cells, with particular attention to the promising role of bacteriophages in this domain. The laboratory's diverse research interests encompass the fields of energy, healthcare, and environmental applications, all revolving around the central theme of utilizing biological processes to engineer nano-scale materials. In this context, the unique characteristics of phages come to the forefront, as their high manipulability makes them indispensable tools for gene modification. Phages are harnessed to cultivate materials for electric batteries, offering enhanced capacity and reaction rates. They find applications in energy storage, conversion, and environmental remediation, making them versatile assets. Beyond these areas, phages also show promise in cancer research, further underscoring their adaptability and potential as platforms for diverse functionalities.

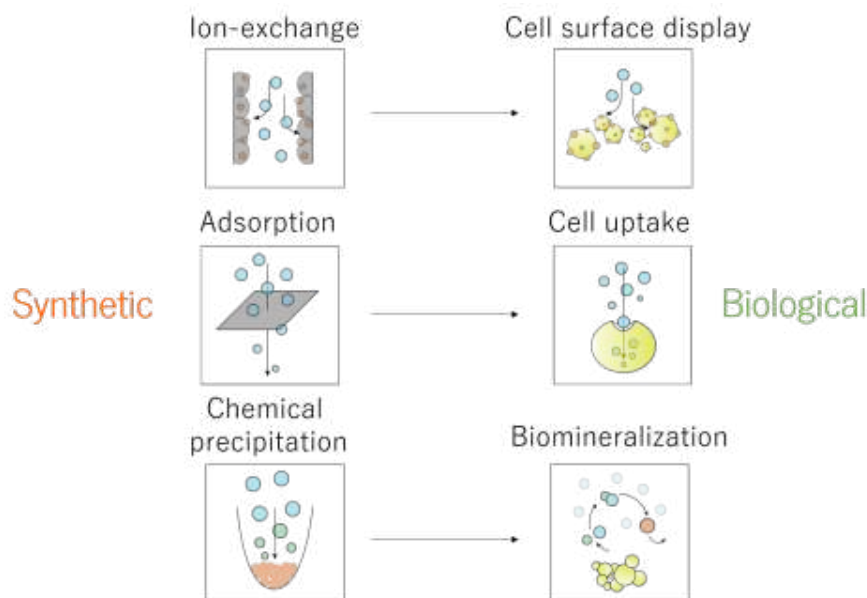


Figure 3. Synthetic vs. Biological metal remediation

Phages and Biomaterials for bioremediation

Dr. Nimrod Heldman | Massachusetts Institute of Technology - USA

Video

This study highlights the versatility of genetic manipulation, focusing on bacteriophages, to transform material characteristics for applications in medicine, energy, imaging, and environmental solutions. Emphasizing the controlled engineering of specific phage components, the research explores real-world applications, including pollution control, climate change mitigation, and viral deactivation. Of particular significance are carbon phage nanofibers that absorb and break down pollutants, providing avenues for protective clothing and virus filtration, notably in the context of the SARS-CoV-2 pandemic. Additionally, the study delves into the realm of hydrogels, using phage modification for light controlled crosslinking to create 3D objects. These hydrogels can integrate therapeutic proteins for potential medical applications, further showcasing the innovative potential of genetic control in material science.

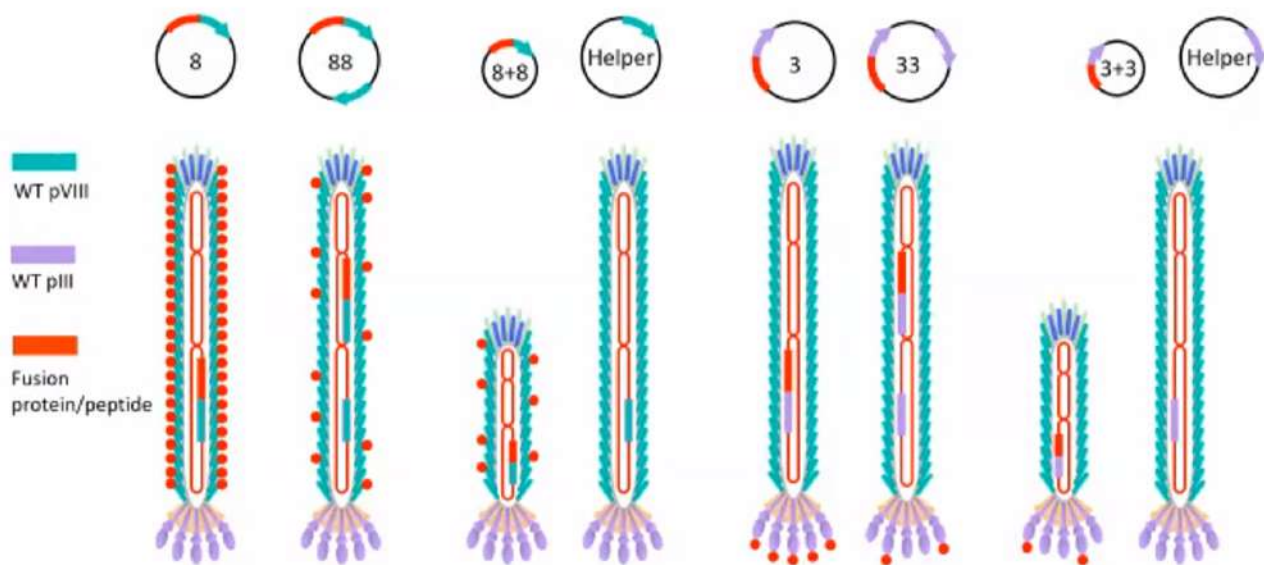


Figure 4. Synthetic vs. Biological metal remediation

Bacteriophages in the environment experiences in Peru

Dr. Miguel Talledo | Universidad Nacional Mayor de San Marcos - Peru

Video

Established at La Chira beach during a 1990s cholera epidemic in Peru, the Basic Characterization Laboratory in Biotechnology for Remediation initially focused on practical *Vibrio* detection using bacteriophage mixtures, with an emphasis on detection rather than comprehensive characterization. Subsequent research led to the isolation of bacteriophages like K14 *Vibrio cholerae*, which showed resistance to chloroform and low infection rates. Phages of *Listeria* (ASCF1) with high temperature and pH tolerance were also isolated, along with other phages like FS01 from *Salmonella typhimurium*. Emphasizing the importance of morphology in characterization, the team also explored phage adsorption on specialized substrates like degradable polyester microfibers loaded with bacteriophages, aiming to standardize substrate applications, with bacteriophages remaining a central focus in their research efforts.

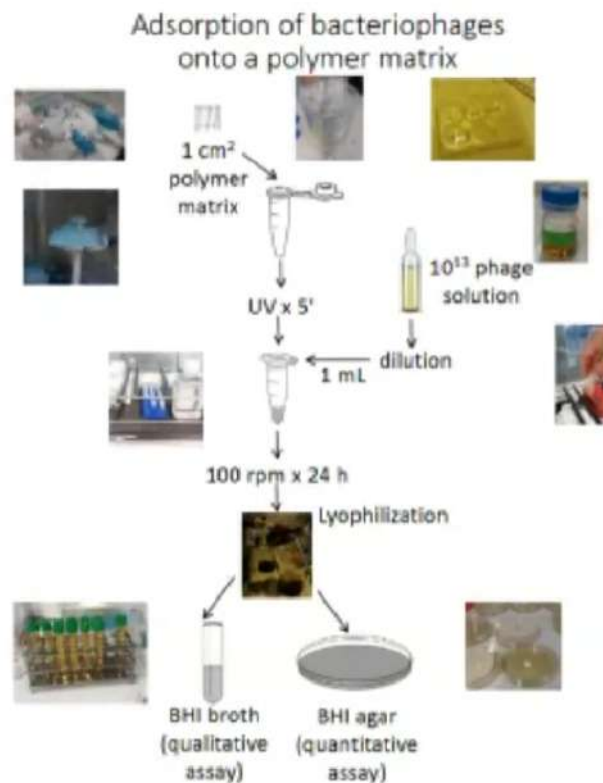


Figure 5. Adsorption of bacteriophages

The Comeback of Phage – Episode II

Dr. Pohl Milón | Universidad Peruana de Ciencias Aplicadas - Peru
Video

Phage therapy, a potential solution for antibiotic resistance, employs phages as carriers for resistance genes. Strategies include early resistance gene detection and novel antibiotic development. Certain phages, like P4, require coexisting phages for replication. Mobile genetic elements play a pivotal role in resistance gene dissemination, prompting environmental impact mitigation efforts. Pandemic-related systems detecting SARS-CoV-2 have been repurposed for resistance gene detection, with a national supply system now established. Collaborative research highlights integron significance in gene transfer. Ongoing research involves river sampling in Oxapampa using mobile detection systems. The incorporation of CRISPR-Cas technology adds specificity to the process, with research exploring its application in natural environments. Undergraduate students actively participate in sample extraction and analysis. Antibiotic development involves conformational change sensor modification and M13 phage-based phage display technology. Combinatorial nucleic acid technologies are also under exploration. Anthropogenic markers signify resistance genes evolving due to human influence, modifying mobile genetic elements.

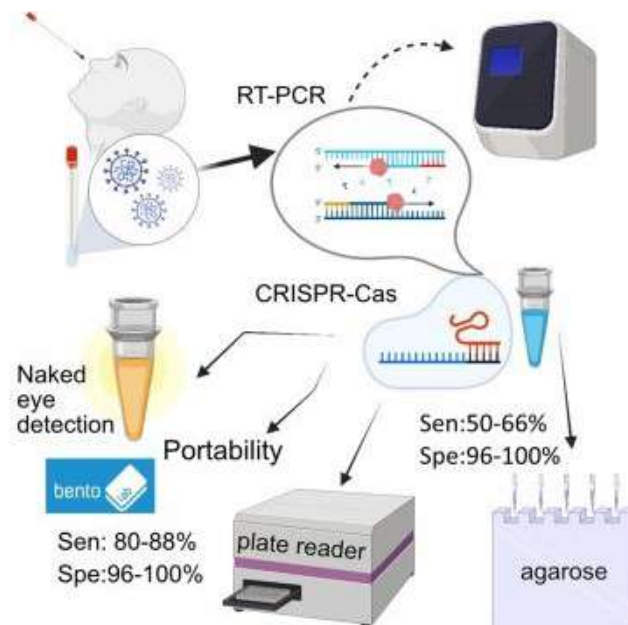


Figure 6. SARS-CoV-2 detection

Bacteriophages and endolysins active against *Staphylococcus aureus*: from the counter to the patient

Dr. Hector Morbidoni | Rosario National University - Argentina

Video

Bacteriophages, abundant biological entities, have been instrumental in advancing biological research and are increasingly harnessed for various applications, including capturing bacteria and biosensing. Argentina has made notable contributions to bacteriophage sequencing. In contrast, endolysins, responsible for bacterial lysis, offer versatility in targeting peptidoglycan domains, making them attractive for therapy. Phages have been isolated from environmental samples, showcasing genetic homology with known phage genes in Latin America. Their potential application in therapy, while showing some variations, is promising. Research on temperate phages facilitates comparative genomics and therapeutic development for diverse strains. Detecting proteins within phage structures is vital for tool development and modification, enhancing their role in proactive research. Organized endolysin domains hold potential for therapeutic development, offering adaptability in recognition and enzymatic activities, shaping the evolving landscape of phage-based solutions.

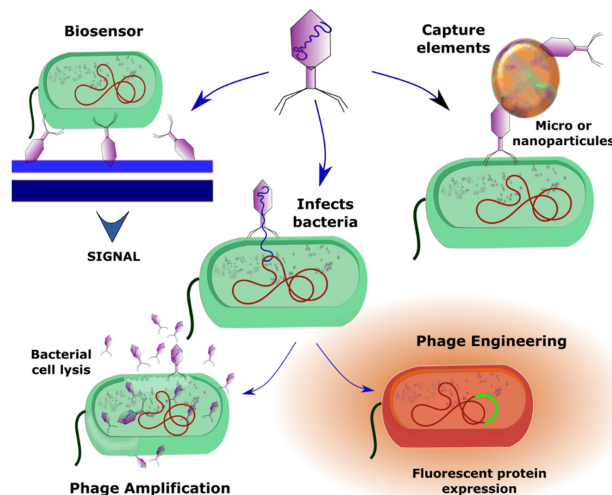
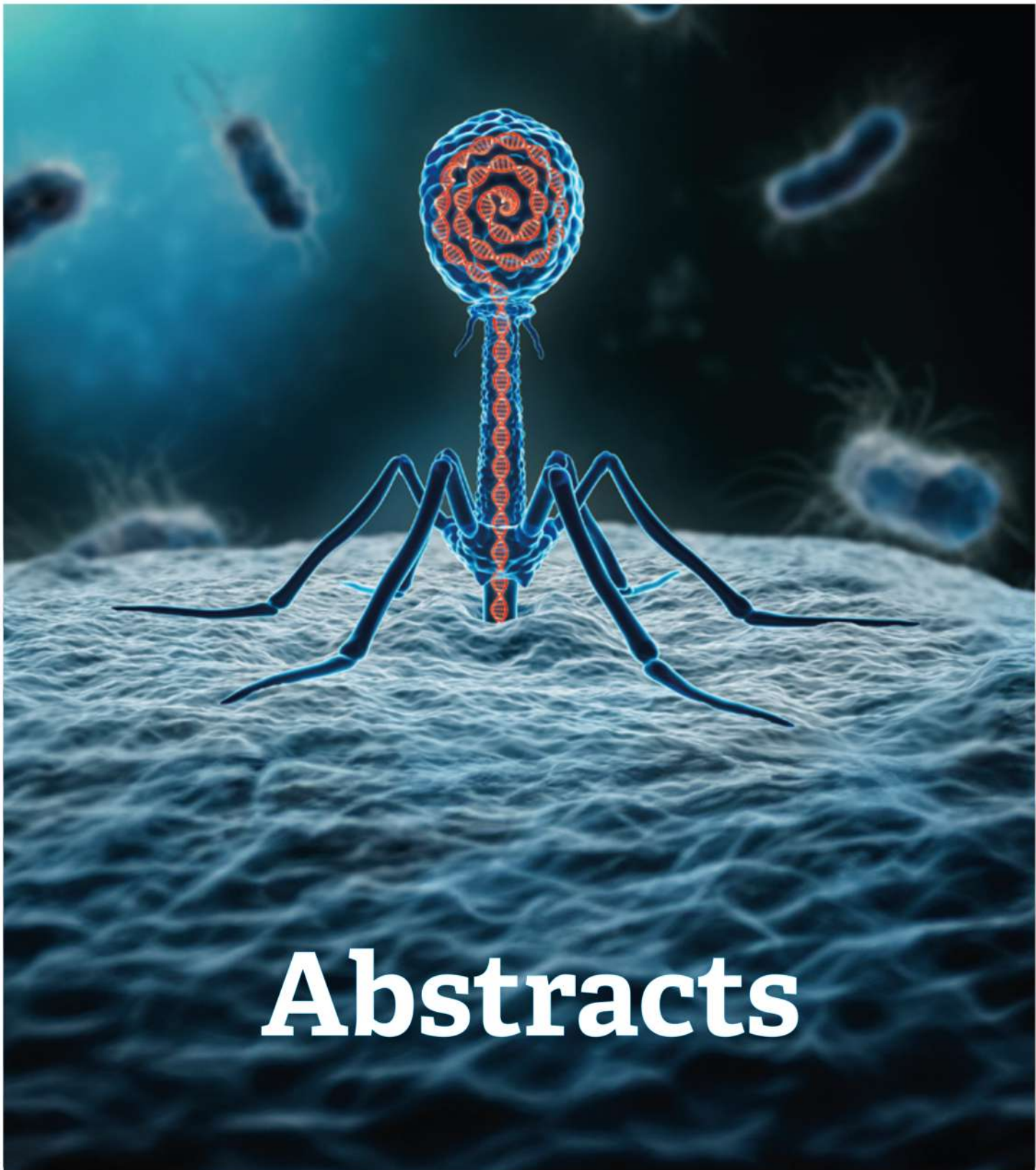



Figure 7. Bacteriophage-based methods for specific bacterial detection




Abstracts

Genetic improvement of bacteriophage $\phi XaF13$ for the biological control of bacterial spot in peppers

Rubén Antonio Olivares-Terrones ¹

Evangelina Esmeralda Quiñones-Aguilar ¹

Guillermo Alejandro Solís-Sánchez ¹

María Inés Siri Tomás ²

Gabriel Rincón-Enríquez ¹


Centro de Investigación y Asistencia en Tecnología y Diseño del Estado de Jalisco¹
Universidad de la República²


Abstract:

Xanthomonas is a genus of Gram-negative bacilli, phytopathogenic in more than 400 species. Bacteriophages are viruses that depend on bacterial cells to propagate. These have increasingly attracted attention for their potential for biological control of diseases caused by *Xanthomonas*. To generate effective biological control strategies, it is necessary to have phages that meet a series of characteristics that allow them to adapt to conditions, so the variety of wild phages may not be sufficient. For this reason, a useful strategy is to improve these phages through random mutagenesis and select those with the best characteristics of resistance to a variety of conditions and biological control. Phage $\phi XaF13$ is a lytic life cycle inovirus with good characteristics for the biological control of bacterial spot in peppers and this work it is aimed to improve it with radiation mutagenesis.

Keywords: *Xanthomonas spp.*, phages, mutagenesis


Engineering p8 protein from *M13 bacteriophage* with high-affinity heavy metal-binding peptides

Jose M. Cisneros Mandujano ¹

Gladys M. Cavero Rozas ¹

Nimrod Heldman ²

Jifa Qi ²

Angela M. Belcher ²

Alberto J. Donayre-Torres ¹

Universidad de Ingeniería y Tecnología¹

Massachusetts Institute of Technology²

Abstract:

This study aims to address heavy metal contamination in water using engineered phage as adsorbent for contaminated water treatment. The objective was to engineer the M13 phage to display along phage surface binding peptides with high affinity to heavy metals. Three peptides with high affinity to Pb(II), Cr(III), and Cd(II) were inserted into the M13 Phage P8 protein gene. The successful genetic editing was confirmed through Sanger sequencing. Transmission electron microscopy (TEM) confirmed structural viability of Pb, Cr, and Cd phages remaining a stable and flexible structure, filamentous shape and a suitable length. TEM revealed phage's structural stability and resilience after Pb(II) exposure and suggested lead adsorption and mineralization along the phage surface. These findings offer promising prospects using engineered M13 phage for heavy metal adsorption from contaminated aqueous solutions. The proposed presents a viable approach for mitigating heavy metal contamination and promoting environmental sustainability.

Keywords: M13 bacteriophage; water remediation; heavy metal ions

Isolation of lytic *Raoultella* bacteriophage

Oscar Nolasco ¹

Mónica Velarde ¹

Ana-Isabel-Flor Gutierrez-Román¹

Universidad Nacional Federico Villarreal¹


Abstract:


Certain species within *Raoultella* and *Klebsiella* genera have become increasingly important due they have been identified as pathogens able to infect plants, some mammals, and humans. As an alternative to antibiotics for bacterial control, several studies have reported the potential use of bacteriophages. The main of this study is to report the isolation of a lytic bacteriophage (*vB_{Rte}*) and its bacterial host BP3, which was identified as a *Raoultella terrigena* strain based on 16S rRNA sequencing. The bacterial host was isolated from a potato with soft rot and the bacteriophage from an agricultural sample using double-layer agar and spot assays. The *vB_{Rte}* was able to inhibit bacterial growth of *Raoultella terrigena* up to MOI 10^{-6} but was unable to infect other bacteria causing soft rot such as *Klebsiella variicola* and *Klebsiella oxytoca* both isolates from banana wet rot disease and *Pectobacterium coratovorum* isolated from potato with soft rot.

Keywords: Bacteriophage; *Raoultella terrigena*; Bacteria


Isolation and characterization of a lytic bacteriophage isolate from chicken intestine and specific for Multidrug-resistant *Salmonella infantis*

Arlene Cuyutupa-Guillen¹

Renzo Punil-Luciano ¹

Egma Mayta-Huatuco ¹

Deisy Ramirez-Soto¹

Enrique Mamani-Zapana ¹


National University of San Marcos¹


Abstract:

Background: The aims were to isolate and characterize a lytic bacteriophage from markets in Lima, Peru, specific for multidrug-resistant *Salmonella infantis*; **Methods:** Phage 52 was isolated by enrichment, processed by spot test and was purified by the double layer method. It was evaluated under different conditions of pH, temperature and exposure to chloroform for 1 hour. It was stored at 25, 4 and -20 °C for 4 months. The optimal multiplicity of infection (MOI) was determined and the one-step growth curve was performed; **Results:** Bacteriophage 52 is specific for *S. infantis* MDR, has an optimal MOI of 0.001, is stable at temperatures from 40 to 60 °C and at pH values from 3 to 10. It showed tolerance to chloroform and has storage stability at temperatures of 4 and -20 °C. Phage 52 has a latency period of 10 minutes and a burst size of 32,93 PFU/cell; **Conclusions:** Phage 52 has great potential to be applied as a control agent for strains for multiresistant *Salmonella infantis* strains.

Keywords: Bacteriophage; Characterization; *Salmonella infantis* MDR

Morphological and molecular characterization of a lytic bacteriophage isolated from soils from bean crops

Erika Camacho Beltran ^{1,2}

Gabriel Rincón Enríquez ¹

Melina López Meyer ²

Evangelina Esmeralda Quiñones Aguilar ¹

Centro de Investigación y Asistencia en Tecnología y Diseño del Estado de Jalisco¹
Instituto Politécnico Nacional²

Abstract:


Given the need to address the biological control of bacterial phytopathogens with a sustainable approach that reduces the use of copper salts and antibiotics, the use of bacteriophages has been considered a strategy in the area of agriculture. Said development depends on the molecular and biological characteristics, therefore, a study of the genome and biology of lytic phages is essential for the development of bacteriophages as biological control agents. The objective was to characterize morphologically and molecularly lytic bacteriophages of bean crops. A bacteriophage of bean bacterial strains from Sinaloa was isolated. Genetic characterization by PCR-RFLPs and morphological characterization by TEM was performed. The lytic bacteriophage ϕ 25-4 measured 37.25 nm from vertex to vertex of the capsid and its genome was single-stranded, circular DNA. This bacteriophage could be used for the biological control of phytopathogenic bacteria that worsen the health of beans in Sinaloa, Mexico.


Keywords: Biological control, phage, *Phaseolus vulgaris*


Isolation of lytic bacteriophages against strains of *Klebsiella pneumoniae* and *Salmonella spp.* multidrug-resistant

Elsa Gladys Aguilar-Ancori ^{1,2}

Duly Nuñez-Carazas¹

Enrique Mamani-Zapana ³

Laura Campo-Pfuyo ¹

Marishani Marin-Carrasco ²

National University of San Antonio Abad¹

University Institute of Tropical Diseases and Biomedicine²

National University of San Marcos³


Abstract:


Background: This study aims to isolate lytic phages against *Klebsiella pneumoniae* and *Salmonella spp.* multidrug-resistant; **Method:** 50 strains of *K. pneumoniae* and 4 multidrug-resistant *Salmonella spp.* strains were characterized as ESBL and carbapenemases, coming from patients of the different hospitals of Cusco. The search for bacteriophages was carried out in the Huatanay River and processed by spot test and double layer. Results: We isolated 14 phages against *K. pneumoniae* and 13 for *Salmonella spp.* obtaining concentrations 3x10⁹ PFU/mL. Likewise, 4 bacteriophages with greater lytic capacity were selected (36% on hospital strains of *K. pneumoniae*) and 03 bacteriophages (80% on *Salmonella spp.* strains). Currently, these data are being processed by transmission electron microscopy for phages, whole genome sequencing using NGS for multi-resistant strains and phages. Conclusion: The lytic phages isolated have potential application in MDR infections against *Klebsiella pneumoniae* and *Salmonella spp.*.

Keywords: Bacteriophage1; *Klebsiella pneumoniae* MDR 2; *Salmonella spp.* MDR 3

Scaling of formulated bacteriophages for biological control of bacterial spot (*Xanthomonas euvesicatoria*) in pepper crops

Alan Hinojosa Román ¹

Evangelina Esmeralda Quiñones Aguilar ¹

Guillermo Alejandro Solís Sánchez ¹

Melchor Arellano Plaza ¹

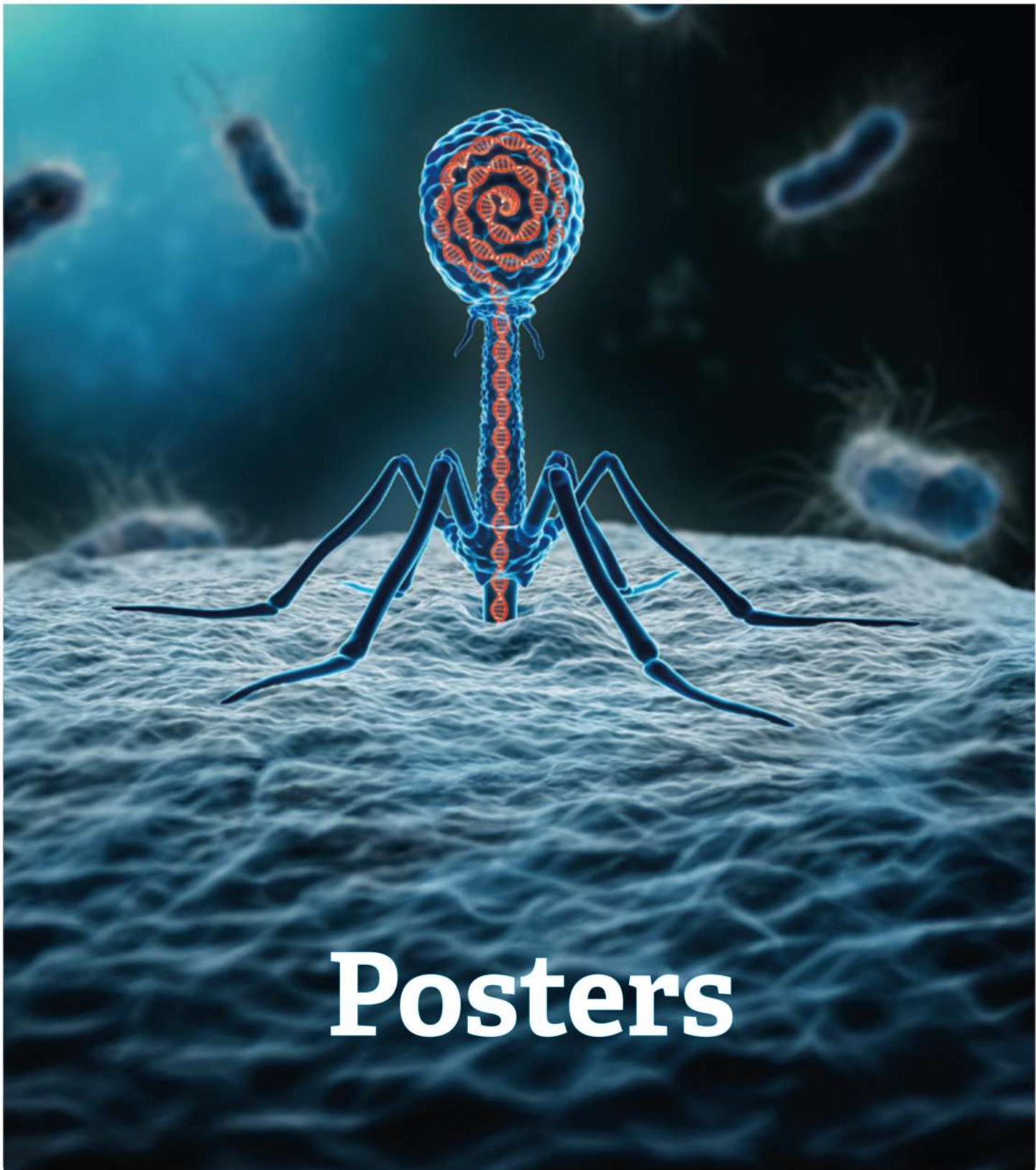
Gabriel Rincón Enríquez ¹

Centro de Investigación y Asistencia en Tecnología y Diseño del Estado de Jalisco¹

Abstract:

The scaling of the photoprotective formulation called Fagolytic based on zinc nanoparticles was evaluated by measuring the growth of its bacteriophages (ϕ XaF13 and ϕ XaF18) in a pilot plant of 20 L and its biological effectivity for *Xanthomonas euvesicatoria* control in pepper crops at greenhouse conditions. The bacteriophages produced in 20 L bioreactors presented concentrations of 1×10^8 UFP/mL y 1.8×10^8 UFP/mL respectively. The statistical analysis was made using the methods of Tukey Pairwise Comparisons and Dunnet Multiple Comparisons with a Control. Both analyses showed that different symptoms behavior were statistically equal in the healthy plants (*control*) and the plants treated with Fagolytic, such as chlorotic spots, necrotic spots, severity scale, and damaged leaf area ($p \leq 0.05$). Given the statistical equality between the Fagolytic treatment, the present commercial preventive treatments and the control in different variables, the formulated bacteriophages show a promising alternative for bacterial spot control in pepper crops.

Keywords: Fagolytic; pepper crops, pilot plant



Posters

Genetic improvement of bacteriophage ϕ XaF13 for the biological control of bacterial spot in peppers.

Rubén Antonio Olivares-Terrones¹, Evangelina Esmeralda Quiñones-Aguilar¹, Guillermo Alejandro Solís-Sánchez¹, María Inés Siri Tomás² and Gabriel Rincón-Enríquez^{1*}

¹Laboratorio de Fitopatología, Unidad de Biotecnología Vegetal, Centro de Investigación y Asistencia en Tecnología y Diseño del Estado de Jalisco, A.C. (CIATEJ) Zapopan, Jalisco, México *grincon@ciatej.mx
²Área Microbiología, Departamento de Biociencias, Facultad de Química, Universidad de la República, Uruguay

GRAPHICAL ABSTRACT

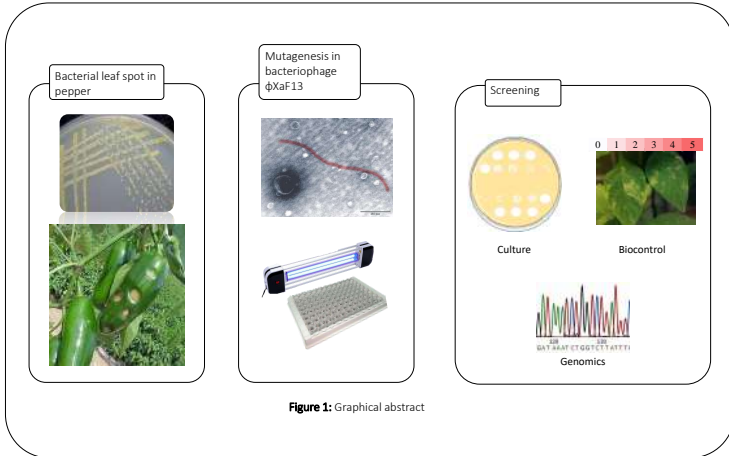


Figure 1: Graphical abstract

INTRODUCTION

Xanthomonas is a genus of Gram-negative bacilli, phytopathogenic in more than 400 species, including pepper (*Capsicum annuum*). Bacterial spot caused by *X. euvesicatoria* pv. *euvesicatoria*, *X. euvesicatoria* pv. *perforans*, *X. hortorum* pv. *gardneri* and *X. vesicatoria* is the bacterial disease that causes the greatest economic damage in pepper crops.

Control of these disease is currently managed in Mexico by cultural and chemical methods, like use of certified seed and copper salts, but there is not commercially available Phage-based product.

Bacteriophages are viruses that depend on bacterial cells to propagate. These have increasingly attracted attention for their potential for biological control of diseases caused by *Xanthomonas* spp. To generate effective biological control strategies, it is necessary to have phages that meet a series of characteristics that allow them to adapt to in field conditions (Ibarra-Rivera, 2019).

Phage ϕ XaF13

Phage ϕ XaF13 is a filamentous bacteriophage of the *inoviridae* family with lytic life cycle, infecting to the phytopathogenic bacteria *X. euvesicatoria*. This phage is highly sensitive to chloroform and have a moderate resistance to UV radiation. Its genome consist of a 7,045-b scDNA encoding 12 ORFs in the positive stand and 2 in the negative strand (Juarez-García, 2022; Solis-Sanchez *et al.*, 2020).

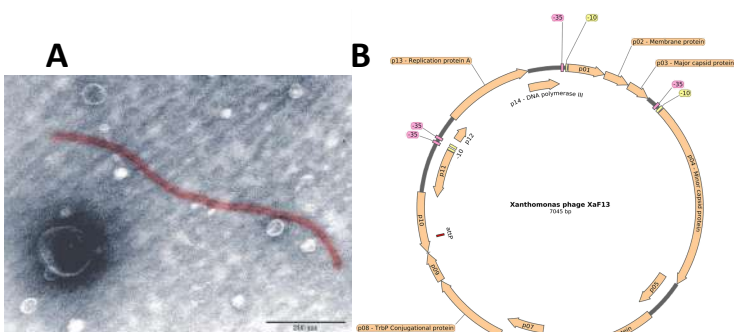


Figure 2. (a) TEM micrograph of the phage ϕ XaF13 (b) ϕ XaF13 genome organization (obtained from Solís *et al.*, 2020)

REFERENCES

Ibarra Rivera, Gabriel. 2019. "Control biológico de la mancha bacteriana (*Xanthomonas axonopodis* pv. *Vesicatoria*) en el cultivo de chile mediante bacteriófagos formulados". Tesis de Maestría, Centro de Investigación y Asistencia en Tecnología y Diseño del Estado de Jalisco, Zapopan.
Juárez García, María Fernanda. 2022. "Genes involucrados en la resistencia a la luz U.V. en bacteriófago ϕ XaF13". Tesis de Maestría, Centro de Investigación y Asistencia en Tecnología y Diseño del Estado de Jalisco, Zapopan.
Solís-Sánchez, Guillermo Alejandro, Evangelina Esmeralda Quiñones-Aguilar, Saul Fraire-Velázquez, Julio Vega-Arreguín, y Gabriel Rincón-Enríquez. 2020. "Complete Genome Sequence of XaF13, a Novel Bacteriophage of *Xanthomonas vesicatoria* from Mexico". *Microbiology Resource Announcements* 9(5). doi: 10.1128/MRA.01371-19.

METHODS

In order to improve the characteristics of phage ϕ XaF13 it will be irradiated with UV-c light to generate random mutations. The stability of the mutant phages will be screened at the in vitro culture level after exposure to various condition to find those more promising.

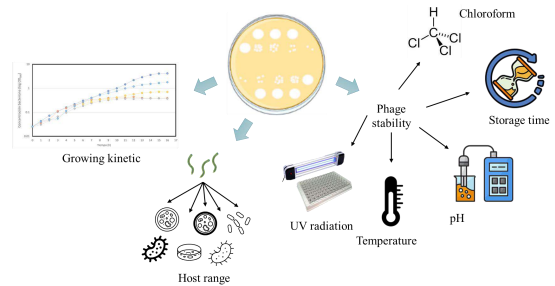


Figure 3. Culture characteristics that will be screened.

In addition to the microbiological culture characteristics, the genomes of the mutant phages will be sequenced to observe the exact sites where gene mutations occur and whether patterns are detected between the appearance of mutations in certain genes and changes in the characteristics of the bacteriophages.

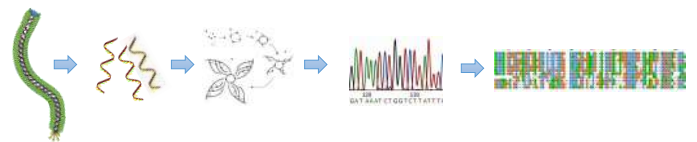


Figure 4. Schematics representation of the genomics analysis.

The ability of mutant phages to control bacterial leaf spot in pepper under greenhouse conditions will also be examined, it will be evaluated according to a severity index, as well as the development of the plant.



Figure 5. Schematics representation of the greenhouse tests.

PERSPECTIVES

The results of this work will help to develop more effective phage-based formulations for the control of bacterial spot in pepper and will also help to understand the mechanism of infection of the ϕ XaF13 phage and of the phages of the *inoviridae* family.

Isolation of lytic Raoultella bacteriophage

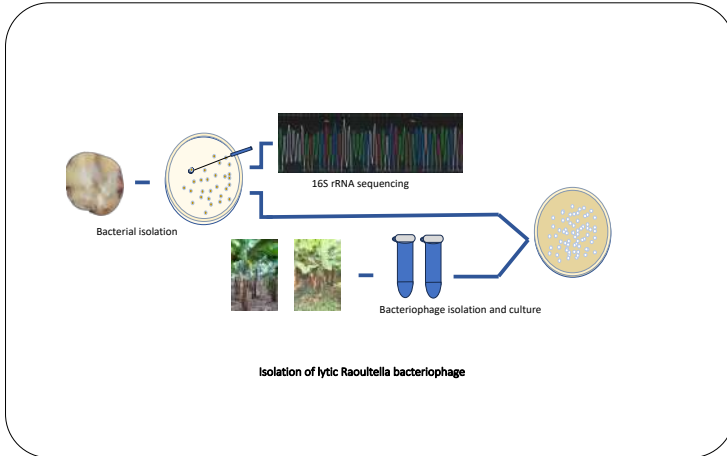
Oscar Nolasco ^{1,2*}, Mónica Velarde ^{1,2} & Ana-Isabel-Flor Gutierrez-Román ^{1,2}

¹ Comunidad de conocimiento GIBBS-UNFV, Facultad de Ciencias Naturales y Matemática, Universidad Nacional Federico Villarreal, Perú.

² Laboratorio de Bioquímica y Biología Molecular, Facultad de Ciencias Naturales y Matemática, Universidad Nacional Federico Villarreal, Perú

* Corresponding author: onolasco@unfv.edu.pe

GRAPHICAL ABSTRACT



RESULTS Y DISCUSSION

The principal bacteria component obtained from potato with soft rot symptoms (BP3) was characterized by microbiology and classified as rods gram-negative and oxidase-negative. Finally, the molecular characterization identified BP3 as *Raoultella terrigena*; the sequence of 1437 nt. of the 16S rRNA gene obtained of BP3 was submitted to the GenBank (Accession number OQ804622). One of thirty different agricultural samples (banana pseudo-stem) displayed a lytic zone. The bacteriophage was amplified and purified by the selection of single clear plaque-forming units (See Figure 1) and was named vB_Rte, using the prefix vB (bacterial virus) and the three-letter host abbreviation Rte (*Raoultella terrigena*). The vB_Rte displayed specificity to *R. terrigena* and was unable to infect phylogenetically close bacteria such as *Klebsiella variicola* and *Klebsiella oxytoca* both isolates from banana pseudostem with symptoms of soft rot. Also was unable to infect *Pectobacterium carotovorum* isolate from potato with soft rot. The stability of vB_Rte was assessed after different storage temperatures and the infectivity remained intact at 12°C storage temperature, while a 10-fold decrease in the number of plaque-forming units was shown after one hour of storage at 55°C or when storage at -20 °C. Null bacteriophage activity was shown after one hour at 65°C. The lytic activity was the same between room temperature to 45°C (See Figure 2). The minimal MOI displayed was 10⁻⁶, with higher dilution non-inhibition to the bacterial growth was shown. The absorbance of the culture broth at different MOIs increased gradually during the first minutes and then decreased according to the decrease in the MOIs value. See Figure 3

INTRODUCTION

Raoultella terrigena is a gram-negative, oxidase negative, re-classified in the Raoultella genus and separated from Klebsiella in 2001 based on 16S rRNA and the rpoβ gene sequences analysis. Microbiological identification of *R. terrigena* is difficult and isolates are frequently misidentified as Klebsiella species [1]. The Raoultella genus are ubiquitous in natural environments (water, soil, plants); *R. terrigena* was isolated from soft rot of chile plants [2]; and among the bacterial consortium for the wheat straw degradation [3]. Likewise, they were reported as opportunistic pathogens in human and animal infections [4,5]. The bacterial role in these biological processes is not clear and efforts should be made to identify and verify its role in vegetable waste degradation and infectious diseases. Over the last few years, some Raoultella bacteriophages have been isolated and reported with the capability to control and balance the bacterial population [6,7]. This short report describes in Peru the isolation of a new phage vB_Rte infecting *Raoultella terrigena*.

METHODS

Bacteria host isolation and identification: The bacteria host was isolated from a potato with soft rot symptoms and subcultured on Tryptic Soy Agar (TSA) plate. The principal bacteria component of the potato with soft rot was identified using the 16S rRNA sequencing.

Lytic bacteriophage isolates: A total of thirty different agricultural samples were assayed for the presence of phages (15 banana pseudo-stem, 1 crop water, and 14 direct soil). The filtrate of each sample was assessed for bacteriophages using double agar overlay and spot test assay.

Host specificity test: The bacterial strains for the specificity test were provided by the "Laboratorio de Investigación de Bioquímica y Biología Molecular - UNFV" (LIBBM-SL10LA105). The bacteriophage infection specificity was tested against *Klebsiella variicola*, *Klebsiella oxytoca*, and *Pectobacterium carotovorum*, all of them related to soft rot. The host specificity test was assessed using double agar overlay and spot test assay.

The Minimum inhibitory multiplicity of infection (miMOI): The miMOI was assessed using fresh culture broth of bacteria host. The bacteriophages were mixed with bacterial culture at different MOIs an incubated at 30°C for 4.5 hours.



Figure 1. Plaques produced by bacteriophage vB_Rte on double-layer solid medium.

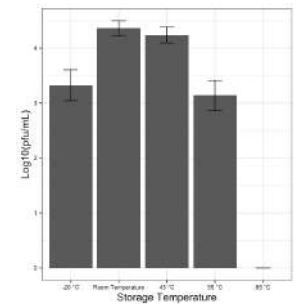


Figure 2. The influence of storage temperature on bacteriophage vB_Rte

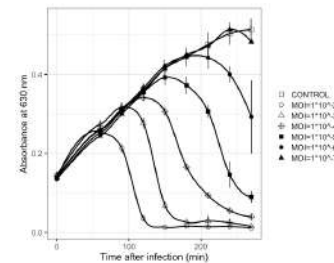


Figure 3. Bacteriophage vB_Rte lysis profile

CONCLUSIONS

This study isolated a lytic bacteriophage specific to *R. terrigena*. The lytic properties showed its potential as an innovative biocontrol agent and specific detection tool for *R. terrigena*. The results of this study need to be expanded for biocontrol as a suitable and realistic antibacterial agent to include in management strategies to the control of degradative bacteria.

REFERENCES

- Dranocourt, M., Bollet, C., Carta, A., et al. Phylogenetic analyses of *Klebsiella* species delineate *Klebsiella* and *Raoultella* gen. Nov., with description of *Raoultella ornithinolytica* comb. Nov., *Raoultella terrigena* comb. Nov. And *Raoultella planticola* comb. Nov. *International Journal of Systematic and Evolutionary Microbiology*, 2001;51(3), 925–932.
- Cubero-Agüero, D., Bressan-Guillén, L., Vidauré-Barahona, D. et al. *Raoultella terrigena* y *Pectobacterium carotovorum* en hortalizas en dos provincias de Costa Rica. *Agronomy: Mesoamericana*, 2021;178–195.
- de Lima Brossi, M. J., Jiménez, D. J., Cortes-Tolalpa, L., et al. Soil-Derived Microbial Consortia Enriched with Different Plant Biomass Reveal Distinct Players Acting in Lignocellulose Degradation. *Microbial Ecology*, 2016;71(3), 616–627.
- Appel, T. M., Quijano-Martínez, N., De La Cadena, E., et al. Microbiological and Clinical Aspects of *Raoultella* spp. *Frontiers in Public Health*, 2021;9. <https://www.frontiersin.org/articles/10.3389/fpubh.2021.686789>
- Wang, Y., Jiang, X., Xu, Z., et al. Identification of *Raoultella terrigena* as a Rare Causative Agent of Subungual Abscess Based on 16S rRNA and Housekeeping Gene Sequencing. *Canadian Journal of Infectious Diseases and Medical Microbiology*, 2016; <https://doi.org/10.1155/2016/3879635>
- Fofanov, M. V., Morozova, V. V., Kozlova, Y. N. et al. *Raoultella* bacteriophage RP180, a new member of the genus Kagunavirus, subfamily Guernevyrinae. *Archives of Virology*, 2019;164(10), 2637–2640.
- Zamani, I., Bouzari, M., Emtiazi, G. et al. Molecular investigation of two novel bacteriophages of a facultative methylotroph, *Raoultella ornithinolytica*: First report of Raoultella phages. *Archives of Virology*, 2019;164(8), 2015–2022.

Scaling of formulated bacteriophages for biological control of bacterial spot (*Xanthomonas euvesicatoria*) in pepper crops.

Hinojosa Román A¹, Rincón Enríquez G¹, Solís Sánchez GA¹, Quiñones Aguilar EE¹ y Arellano Plaza M²

¹Laboratorio de Fitopatología Vegetal, Biotecnología Vegetal, Centro de Investigación y Asistencia Tecnológica y Diseño del Estado de Jalisco, A. C. (CIATEJ). Zapopan, Jalisco., El Bajío, Tel: 33 3345 5200, CP. 45019.

²Planta piloto, Biotecnología Industrial, Centro de Investigación y Asistencia en Tecnología y Diseño del Estado de Jalisco, A. C. (CIATEJ). Zapopan, Jalisco., El Bajío, Tel: 33 3345 5200, CP. 45019.

GRAPHICAL ABSTRACT

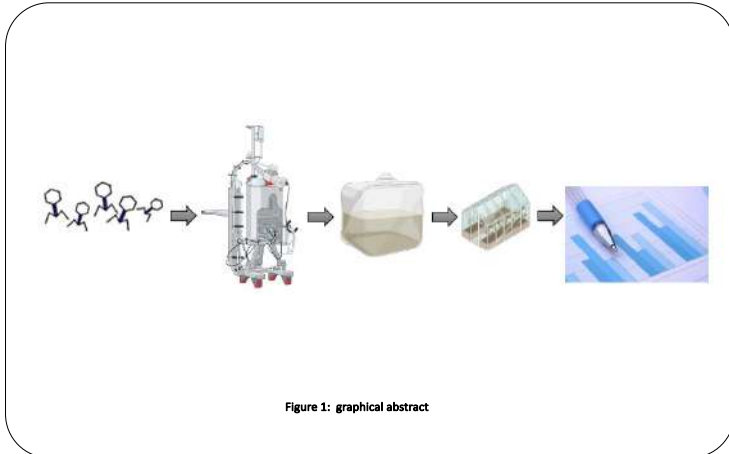


Figure 1: graphical abstract

INTRODUCTION

The current methods in Mexico to control the *Xanthomonas euvesicatoria* in pepper crops are primarily based on prevention, as chemical control proves intricate due to issues such as the bacterium's rapid resistance development against active compounds like copper-based compounds and antibiotics (Ibarra, 2019). Consequently, these classical pest control methods lead to long-term repercussions on pepper cultivation, elevating production costs, diminishing effectiveness, and disrupting the balance of the natural ecosystem.

Given the problems arising from ineffective strategies in controlling *Xanthomonas* spp. pests, interest has surged in the utilization of biocontrol through viruses known as bacteriophages, which have coexisted with bacteria and crops in natural conditions (Nakayinga *et al.*, 2021). When applying a bacteriophage in natural or greenhouse conditions to combat the *X. euvesicatoria* in *Capsicum annuum* agricultural production, one must consider challenges that may arise and prove challenging to manage, such as UV radiation, high temperatures, and the bacterium's potential immunity generation. UV radiation stands as a primary limitation to bacteriophage effectiveness in pest control, central to the Fagolytic product/invention.

In this study, an upscaling of Fagolytic was pursued, composed of zinc nanoparticles and the lytic phages ΦXaF13 and ΦXaF18, to a pilot plant scale of 20 L. Its biological effectiveness in controlling *X. euvesicatoria* in pepper crops under greenhouse conditions was evaluated.

METHODS

Bacteriophages were produced at the pilot plant level in a 20 L Applikon bioreactor. Centrifugation and filtration of the product was carried out using a Thermo Scientific Centrifuge Multifuge 3XR and a Millipore Tangential Filter. Finally, the biological effectiveness of Fagolytic was evaluated in pepper crops established in a plastic greenhouse using the methods of Tukey Pairwise Comparisons and Dunnett Multiple Comparisons ($p \leq 0.05$) with a control employing chlorotic spots, necrotic spots, number of leaves with symptoms, defoliation, severity scale and damaged leaf area per leaf as variables.

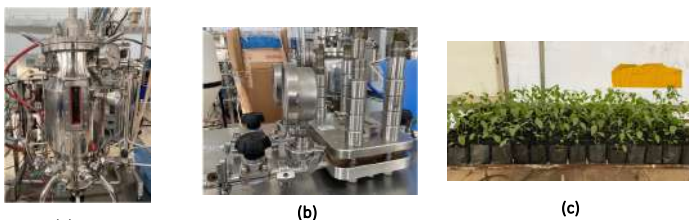


Figure 2. (a) Bioreactor (b) Tangential Filter (c) Pepper plants in greenhouse

RESULTS Y DISCUSSION

The results obtained from the titrations of batches 1 and 2 of bacteriophages produced in the 20 L bioreactors (Table 1) showed suitable concentrations for the biological effectiveness test in pepper crops under greenhouse conditions that was carried out at the end of the formulation of Fagolytic products. The concentration dimensions of the phages are close to those obtained by Payan (2022).

The statistical analysis made by the methods of Tukey Pairwise Comparisons and Dunnett Multiple Comparisons ($p \leq 0.05$) with a control showed that in many of the symptoms provoked by the *X. euvesicatoria* bacterial spot disease, such as chlorotic spots, necrotic spots, severity scale, and damaged leaf area has statistically equal behavior when is treated with Fagolytic as when is with the present commercial preventive treatments, and in a few cases with the control treatment (healthy).

Table 1. Bacteriophage batches

Batch (date of production)	BACTERIOPHAGE	CONCENTRATION (UFP/mL)
1 (2nd of march)	ΦXaF13	1x10 ⁸
2 (13th de march)	ΦXaF18	1.8x10 ⁸

Chlorotic spot (Day 21)			
Tukey Pairwise Comparisons ($p \leq 0.05$)			
Factor	N	Mean	Grouping
Serenade	19	31.58	A
Formulated phages	19	28.58	A
Non-formulated phages	19	26.68	A
Sick plant	19	26.05	A
Kasumin	19	23.84	A
Fagolytic	19	18.68	A B
Healthy plant	19	1.211	B

Figure 3. Pairwise comparisons of chlorotic spots provoked by bacterial spot disease.

Necrotic spot (Day 21)			
Tukey Pairwise Comparisons ($p \leq 0.05$)			
Factor	N	Mean	Grouping
Serenade	37	18.95	A
Formulated phages	37	18.11	A
Non-formulated phages	37	17.8	A
Sick plant	37	16.92	A
Kasumin	37	15.24	A
Fagolytic	37	11.54	A B
Healthy plant	37	1.216	B

Figure 4. Pairwise comparisons of necrotic spots provoked by bacterial spot disease.

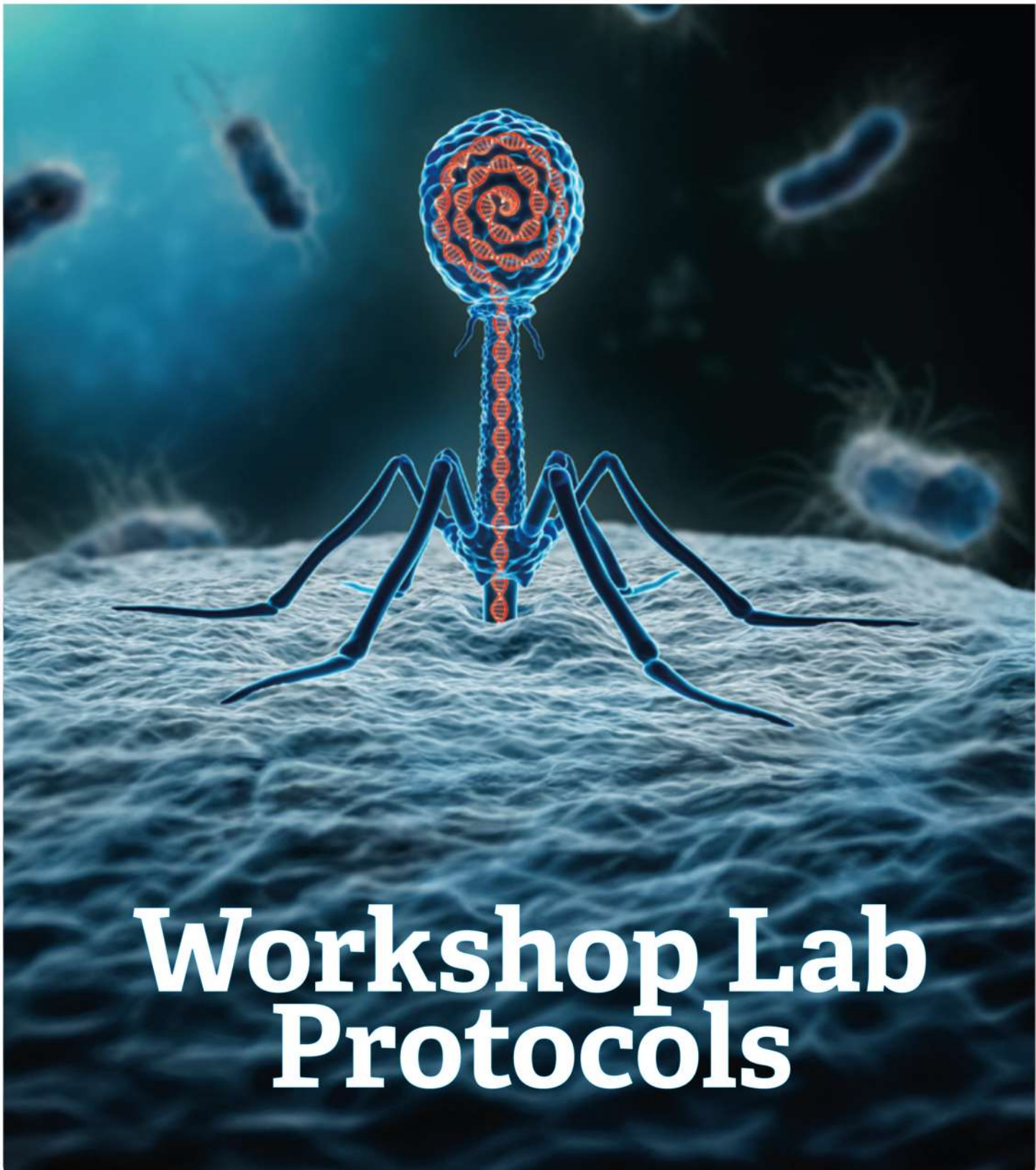
Given the present results, the Fagolytic treatment in pepper crops at greenhouse conditions shows statistically the same behavior in symptoms caused by the bacterial spot disease of *X. euvesicatoria* when the culture is healthy as when is treated previous the infection.

CONCLUSIONS

Given the statistical equality between the Fagolytic treatment, the present commercial preventive treatments and the control in different variables, the formulated bacteriophages show a promising alternative for bacterial spot control in pepper crops.

REFERENCES

- Ibarra Rivera, «Control biológico de la mancha bacteriana (*Xanthomonas axonopodis* pv. *vesicatoria*) en el cultivo de chile mediante bacteriófagos formulados.» Centro de Investigación y Asistencia en Tecnología y Diseño del Estado de Jalisco, A. C., Zapopan, 2019.
- Nath L., 2008. Colorimetric method for determination of sugars and related substances. *Anal. Chem.* 20, 350-356.
- J. D. Payan Almanza, «Formulación a base de bacteriófagos para el biocontrol de la mancha bacteriana en el cultivo de solanáceas: producción en planta piloto y evaluación pre-comercial.» Centro de Investigación y Asistencia en Tecnología y Diseño del Estado de Jalisco, A. C., Zapopan, 2023.
- R. Nakayinga, A. Makumi y V. Tumuhaise, «*Xanthomonas* bacteriophages: a review of their biology and biocontrol applications in agriculture.» *BMC Microbiol.* vol. 21, p. 219, 2021.



Workshop Lab Protocols

Workshop 1

Using of Molecular Cloning Techniques for phage genome (vector M14KE) editing

Video

Experience 1

Gibson assembly and primer design

In 2009, Dr. Daniel Gibson and his colleagues first described the Gibson assembly technique. Gibson assembly is a single reaction technique for the assembly of multiple overlapping DNA fragments. It enables the insertion of an insert into the desired vector. The process begins with DNA fragments, both insert and open vector, possessing regions of homology at their ends, typically generated through PCR. DNA fragments are then incubated together with an enzyme master mix comprising three different enzymes:

1. An exonuclease, which trims back the 5' ends of the fragments, creating long overhangs that allow the single-stranded regions with homology to anneal.
2. A polymerase, which fills in the gaps.
3. A DNA ligase, which seals the nicks of the annealed and filled-in gaps.

One of the remarkable aspects of this enzyme mix is that they all function at the same temperature. Consequently, the entire reaction can be completed in about an hour at 50°C or lower. After this short incubation period, the sample is ready for immediate transformation into competent cells.

Materials and equipment:

- Snapgene

Sanger sequencing analysis

Automated DNA Sequencers produce a chromatogram with four colors, representing the results of the sequencing process, along with a text file containing the sequence data as interpreted by a computer program. The characteristics of Sequencing Chromatograms are as follows.

Workshop 2

Amplification of recombinant phages at high scale. Purification using centrifugation and chemical precipitation

Video

Experience 1: Plasmid DNA quantification in Nanodrop

Reagents:

- 1.5 mL Eppendorf Tubes with plasmid
- 1.5 mL Eppendorf Tubes with blank (water or buffer)

Materials and equipment:

- UV/Vis Spectrophotometer
- Tips (10 ul)

Protocol: Plasmid DNA quantification in Nanodrop

1. Add 2uL of Millipore deionized water to initialize the instrument.
2. Once completed, remove the fluid with Kimwipe and add 2uL of the blank solution and press "Blank". This will use the absorbance spectra of the solution as a reference for the sample measurement.
3. Place 2uL of the plasmid DNA solution on the Nanodrop pedestal and run the machine.
4. Record the absorbance.

Experience 2: Bacterial transformation

Description:

This manual provides a step-by-step protocol for the transformation of XL1-Blue competent cells using the M13KE gIII Cloning Vector. The transformation process involves the introduction of the M13KE vector into the bacterial cells, allowing for the expression of desired genes or DNA sequences.

Reagents:

- SOC medium
- XL1-Blue Competent Cells Catalog #200249
- M13KE gIII Cloning Vector (10 ng/ul)

Materials and equipment:

- 1.5 mL Eppendorf Tubes
- 15 mL tubes
- Tips (10, 200 and 1000 ul)
- Water bath at 42 °C
- Ice bucket filled with ice
- Incubator at 37 °C

Transformation Protocol

1. Thaw one vial of chemically competent cells on ice and transfer 100 ul from this tube to a 1.5 ml eppendorf tube. Work in ice.
2. Add 5 ul of M13KE vector (10 ng/ul) to the tube to the competent cells (You are adding 50 ng of vector). Mix gently by pipetting up and down or by flicking the tube 4–5 times. Do not vortex.
3. Place the mixture on ice for 30 minutes. Do not mix.
4. Heat shock at 42°C for 30 seconds. Do not mix.
5. Transfer tube to ice for 2 minutes.
6. Add 900 µl of room-temperature SOC media to the tube.
7. Incubate the bacteria transformation reaction at 37°C for 60 minutes. Shake vigorously (250 rpm) or rotate.

Experience 3-a: Tetracycline Preparation

Reagents:

- Tetracycline hydrochloride
- 70% ethanol

Materials and equipment:

- One tube 15 ml
- Ice bucket filled with ice
- Laminar Flow hood

Protocol Blue LB agar

1. To prepare 10 ml of 20 mg/ml tetracycline stock solution, weigh out 200 mg of tetracycline. Transfer it to a 15 ml screw-capped tube. Add 9 ml of 70% ethanol. Mix until all tetracycline dissolves completely. Once the content is mixed, make up the volume to 10 ml.
2. Store at -20°C for a long time. Protect the solution from light, 0.22 μ m syringe filter.

Experience 3-b: Blue agar preparation

Reagents:

- LB medium (Dehydrated powder)
- Bacto Agar in powder
- X-Gal solution (50 mg/ml) (Promega)
- Tet 1000X
- IPTG 100mM (100X)
- Milli-Q Water

Materials and equipment:

- ~20 units of Petri Dish Boxes
- Tips (1000 ul)

Protocol:

1. To prepare 500 ml of LB-agar, in 500 ml of sterile water add:
 - 5 g/L Bacto-Tryptone
 - 2.5 g/L yeast extract
 - 2.5 g NaCl
 - 7.5 g/L Agar.
2. Autoclave and cool.
3. Then add IPTG, Xgal, and Tet solutions.
 - Add 0.4 ml of X-Gal solution (50mg/ml) (*Promega*).
 - Add 5000 ul of IPTG (100mM).
 - Add 0.5 mL Tet 1000X.
4. Mix before using. Pour into Petri dishes (~25 mL per plate).

Workshop 3

Transformation of phage genome into XL1-Blue cells using chemical transformation

Video

Experience 1: Phage Plating and aliquoting with Top agar

Description:

This manual provides a detailed protocol for conducting phage titering, a technique used to determine the concentration or titer of bacteriophages present in a sample. The titering process involves serial dilutions of a genetically transformed bacterial culture and subsequent plating on agar plates to visualize the formation of phage plaques. **Reagents:**

- LB Agar Plaques with IPTG/XGal/Tet
- Top Agar
- LB Medium
- 4 mL Overnight Culture of XL1-Blue Competent Cells
- Solution with Phage

Materials and equipment:

- 1.5 mL Eppendorf Tubes
- 15 mL tubes
- 10 mL Pipette
- Micropipette Tips
- Incubator at 37 °C
- Microwave

Protocol

1. Microwave Agarose Top and store at 50-60°C. High temperature can kill samples.
2. Place 4 plates in the incubator at 37 °C upside down.

3. Do 10 serial dilutions ($1 : 10$ to $1 : 10^{10}$) with the phage solution tube. In ten 1.5 mL eppendorf tubes, add 90 μ L LB medium.
 - (a) **Dilution tube 1 ($1:10$):** Take 10 μ L of phage solution tube and resuspend it in the 90 μ L dilution media. Mix by pipetting.
 - (b) **Dilution tube 2 ($1 : 10^2$):** Add 10 μ L of Dilution tube 1 to the tube with 90 μ L LB medium.
 - (c) **Dilution tube 3 ($1 : 10^3$):** Add 10 μ L of Dilution tube 2 to the tube with 90 μ L LB medium.
 - (d) Repeat the process until you reach Dilution tube 10 ($1 : 10^{10}$).
4. Aliquot 100 μ L Overnight Culture of XL1-Blue Competent Cells in eleven Eppendorf tubes and label them as Tube 0, Tube 1, Tube 2, Tube 3 until Tube 10.
5. Add 10 μ L of the phage solution tube to Tube 0, pipetting to mix it. Add 10 μ L of each dilution tube to its respective tube that has 100 μ L E. coli overnight culture, pipetting to mix it.
6. Retrieve plates and Agar Top. Label the back of the plates as Plate 0, Plate 1, Plate 2, and Plate 3.
7. Aliquot 3mL Agar Top into 15 mL tubes.
8. Pipette the 110 μ L Phage/E. coli solution (from Tube 0, Tube 1, Tube 2, Tube -3, and so on) into the 3mL Agar top and mix. You can use a vortex on low or swirl it by hand.
9. Pour the newly mixed solution onto a correspondingly labeled petri dish, swirling it to evenly cover the full area.
10. Repeat this process for each of the dilutions and Agar top samples.
11. Once plates have cooled and solidified, place them back in the incubator at 37 °C upside down overnight.

Note: The Dilution tube 8 ($1 : 10^8$) showed a good distribution of colonies infected with phage. For the workshop, the Phage Plating will be using Dilution tube 8 ($1 : 10^8$). We will start from the 4th step of this protocol.

Workshop 4

Titering of M13 bacteriophages for selection of cells expressing phages Quantification of phage concentration using a spectrophotometer

Video

Experience 1: Large scale amplification of bacteriophage

Description: This manual provides a detailed protocol about the amplification process of M13KE bacteriophages. This step involves infecting a bacterial culture with a small amount of bacteriophage so that it can replicate a larger population of new phages. Amplification can be performed in multiple rounds to increase the quantity of bacteriophages.

Reagents:

- Millipore DI water
- LB (Luria-Bertani) media
- $MgCl_2 \cdot 6H_2O$
- Tetracycline (TET)
- E. coli colony (XL-1 Blue)
- Stock solution of roughly $1e13$ virions/mL phage
- 10% bleach solution

Materials and equipment:

- Erlenmeyer Flasks (500mL)
- Autoclaved 250 mL Erlenmeyer flask
- 2L flask
- Polycarbonate centrifuge tubes
- Centrifuge with JLA 81,000 rotor
- 4°C refrigerator

- NanoDrop UV/Vis spectrophotometer
- Pipettes and tips
- Vortex mixer
- Microcentrifuge
- 40mL centrifuge tube

Amplification and purification Protocol

Day 0: Materials preparation and 4mL E. coli OC

1. Combine 500 mL Millipore-DI water with 10.5 g LB (in powder) in a large 1L flask and 0.5 g of MgCl₂·6H₂O. Prepare 2 Erlenmeyer Flasks. Autoclave the flasks with the solution inside. Cool the LB solution below 45°C and add 0.5mL of Tetracycline solution to each flask.
2. Combine 25 mL Millipore-DI water with 0.515 g LB (in powder) in a small 100 mL flask. Prepare 2 Erlenmeyer Flasks. Autoclave the flasks with the solution inside. Cool the LB solution below 45°C and add 25 ul of Tetracycline solution to each flask.
3. In a 15 mL tube, add 4mL LB, 4 ul of Tet 1000X, and a sample of E. coli X11-Blue. Make sure there is 167mL of PEG/NaCl for every liter of LB being amplified. If not, add 150g of PEG-8000 and 110g NaCl to 750mL of Millipore DI H₂O.

Day 1: Prepare Overnight Culture (OC)

1. Small flask with E.coli: Add 250 ul of E. coli overnight culture with 25uL of TET to the flask with 25 mL of LB.
2. Small flask with Phage: Add phage stock, 250 ul of E. coli overnight culture with 25uL of TET to the flask with 25 mL of LB. Incubate overnight in a 37°C incubation shaker.

Day 2: 24 Hour Virus Amplification

1. Add 10 mL of Small flask with E.coli to 1L flask (prepared in Step 1).
2. Measure optical density at 600 nm.
3. When optical density is 0.4, add 10 mL of Small flask with Phage to 1L flask.
4. Incubate for 24 hours by shaking at 37°C and 225-250 rpm (shake for at least 6-7 hours).

Experience 2: Purification using centrifugation and protein precipitation

Reagents:

- Millipore DI water
- PEG-8000
- NaCl
- 1x PBS solution

Materials and equipment:

- Autoclaved 250 mL Erlenmeyer flask
- 15 mL centrifuge tubes
- 50 mL centrifuge tubes
- Centrifuge for 15 mL tubes
- Centrifuge for 50 mL tubes
- 4°C refrigerator
- NanoDrop UV/Vis spectrophotometer
- Pipettes and tips
- Vortex mixer

Day 3: Phage Separation and Purification Round 1

1. Extract the flask from the shaker and transfer the contents of each flask into separate polycarbonate centrifuge tubes.
2. Perform centrifugation on the overnight culture using the largest rotor available at a speed of 8,000 rpm for a duration of 30 minutes. This procedure should lead to the formation of a bacterial pellet at the bottom of the centrifuge tube. Meanwhile, the virus, due to its significantly smaller mass compared to bacterial cells, will remain suspended in the supernatant.
3. Once the centrifugation cycle is completed, decant the supernatant into a flask with a capacity of 1L. Add PEG/NaCl to each in a volume that corresponds to 1/5th of the volume of the virus solution.

4. Store the flask in a refrigerator set at a temperature of 4°C overnight. The addition of PEG and salt serves to draw water molecules away from the virus, inducing its separation from the fluid. Over the span of several hours, the virus will initiate a process of aggregation and precipitation, causing it to settle out of the solution.
5. Utilize a disposable spatula to scrape off a segment of the bacterial pellet and preserve it in a 1.5mL round bottom centrifuge tube. This sample will be reserved for DNA extraction purposes. Store this tube in the refrigerator set at a temperature of 4°C.

Day 4: Phage Purification Round 2

1. Transfer solution from Step 16 into centrifuge tubes. Spin at a speed of 8,000 rpm for 60 minutes. Make sure not to shake the solution too much.
2. Once the spinning is done, you should see a white pellet at the bottom of the tube. Quickly pour out and get rid of the liquid part into a big bottle with around 20mL of a liquid that's 10
3. Dissolve the pellet by putting 10mL of PBS (1X) in each tube. To do this, gently shake the bottles and move a pipette up and down close to the pellet. Collect the phage solution, which is the liquid part, in a 50 mL centrifuge tube. Centrifuge these tubes at 14000 rpm for 30 minutes. Transfer supernatant into clean 50 centrifuge tubes. Discard pellets (residual impurities).
4. Add 2mL of PEG/NaCl solution into each of the tubes. Put the tubes from Step 22 into a refrigerator around 4°C. Leave them in there overnight.

Day 5: Final Phage Purification and Collection

1. Spin the phage solution in the centrifuge at a speed of 14,000 rpm for 45 minutes.
2. Completely mix the solid part (pellet) in each tube with 2mL of PBS (1x).
3. Move the solution into 1.5 Eppendorf tubes. Spin them at a speed of 14000 rpm for 10 minutes.
4. Move the supernatant to new clean 1.5mL eppendorf tubes.
5. This is now your final phage solution.

Day 5: Quantification with the NanoDrop

1. Choose "UV/Vis" on the main menu.

2. Put 2uL of Millipore DI water to start the device.
3. When that's done, clean away any liquid with a Kimwipe cloth, and then add 2uL of PBS (1x) solution.
4. Push the "Blank" button. Put 2uL of phage solution from Step 27 onto the platform of the Nanodrop.
5. Write down the absorbance numbers at 269 and 320 nm.

Use the following formula to find out the final phage concentration:

$$[M13] = \frac{(A_{269} - A_{320}) \cdot 6 \times 10^{17}}{7225}$$

