Lytic phages displayed protective effects against soft rot-causing

***Pectobacterium* sp.**

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Abstract:
Soft rot caused by *Pectobacterium* sp. is responsible for significant losses in vegetable
production worldwide. Methods for the effective control of soft rot are limited and are
primarily based on good agricultural practices. The use of bacteriophages as biocontrol
agents appears to be a promising alternative to combat phytopathogens. In this study, we
investigated the efficacy of lytic phages recovered from symptomatic tissues and
environmental samples against soft rot caused by *Pectobacterium carotovorum* subsp.
carotovorum. Three bacteriophage isolates, designated as vB_PcaP-A3, vB_PcaM-D1, and
vB_PcaM-J3, were observed to effectively lyse *P. carotovorum* subsp. *carotovorum*. Phage
vB_PcaP-A3 exhibited virion morphology similar to the members of the podovirus group,
while phages vB_PcaM-D1 and vB_PcaM-J3 showed myovirus morphology based on
transmission electron microscopy. The optimal multiplicity of infection (MOI) differed
greatly among the three phages. All three phages survived incubations at 30°C, 40°C and
50°C and pH conditions ranging from 3.0 to 9.0, but were all inactivated at 60°C and at pH
12. Both monophage and cocktail preparations were effective in inhibiting the growth of *P.*
carotovorum subsp. *carotovorum* in the *in vitro* challenge tests. In the semi-*in planta* assays,
while treatment with cocktail preparations completely inhibited the development of soft rot in
tissue slices, monophage treatments not only resulted in significant reduction of tissue
maceration in slices, but also showed protective effect against soft rot in tubers. Overall,
these results demonstrate the efficacy of phages vB_PcaP-A3, vB_PcaM-D1, and vB_PcaM-
J3 for the biocontrol of soft rot caused by *P. carotovorum* subsp. *carotovorum.*

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1. Introduction

Bacterial soft rot disease causes serious damage among vegetable and fruit crops worldwide (Bhat et al., 2010). Annual post-harvest losses due to bacterial soft rot vary from 15-30% of the harvested crops and at least 30-50% of the 65 million tons of vegetables in public markets (Bhat et al., 2010; Monjil et al., 2021; Rahman et al., 2012). Soft rot disease results from infections by several types of bacteria, but most commonly by members of the soft rot *Pectobacteriaceae* (SRP) including *Pectobacterium carotovorum* subsp. *carotovorum*, *P. aroidearium*, *P. atrosepticum*, *P. brasiliense*, *P. betavasculorum* and *P. wasabiae,* which were identified to cause soft rot both in storage and in transit (Adeolu et al., 2016; Agyemang et al., 2020; Charkowski, 2018; Czajkowski, 2015; Gardan et al., 2003; Nabhan et al., 2013; Pérombelon, 2002; Portier et al., 2019; Wallace, 1948). *Pectobacterium* species have the broadest host range among soft rot bacteria, and are considered as the leading cause of soft rot disease in various vegetable crops (Davidsson et al., 2013; Lim et al., 2017). During infection, SRPs produce enzymes like pectate lyase, cellulase and polygalacturonase, which are capable of breaking down the plant cell wall leading to plant tissue maceration and decay (Charkowski et al., 2012; Li et al., 2019; Mattinen et al., 2007; Muturi et al., 2019). There is still limited information on the predominant SRPs affecting vegetable crops in the Philippines. A previous report by Daengsubha and Quimio (1980) identified *P. carotovorum* subsp. *carotovorum* as the major cause of bacterial soft rot affecting crucifers, carrot, and tomato.

Methods used to control soft rot disease include cutting-off and disposal of infected tissues during or after harvest, wrapping with paper, treatment with hot water, and the use of bactericides, such as alum powder (aluminum potassium sulfate), formaldehyde, copper sulfate, and sodium or calcium hypochlorite (Bayogan et al., 2011; Bhat et al., 2010;

Czajkowski et al., 2011; Liao et al., 2003). However, these methods only slow down the

development of soft rot, and the use of chemical bactericides were reported to be associated

with the detection of copper- and antibiotic-resistant phytopathogenic bacterial strains

(Behlau et al., 2011; Cooksey, 1990; McManus et al., 2002; Petriccione et al., 2017; Roach et

al., 2020; Sundin & Wang, 2018). Due to the limited success of the abovementioned control

methods, and the currently growing global threat of antimicrobial resistance, there has been

an increased interest in the use of bacteriophages or phages (viruses of bacteria) as biocontrol

agents against pathogenic bacteria (Buttimer et al., 2017; Frampton et al., 2012; Jones et al.,

2007; Jones et al., 2012).

Bacteriophages (phages) are viruses that infect and lyse bacterial cells. When environmental

conditions are favorable for infection, phages can effectively reduce bacterial populations, as

in the case of lytic phages that undergo rapid replication (Hadas et al., 1997; Sulakvelidze et

al., 2001). Since phages have high host-specificity and have no direct negative effects on

non-host cells, these allow the use of phage therapy to be a promising biocontrol method

against soft rot caused by *Pectobacterium* sp. (Buttimer et al., 2017; Lim et al., 2017). The

first *Pectobacterium* phage that was sequenced, PP1, was reported in 2012 and several

additional novel phage isolates were subsequently reported (Buttimer, Lucid, et al., 2018;

Buttimer et al., 2020; Kalischuk et al., 2015; D. H. Lee et al., 2012; Lim et al., 2017; Lim et

al., 2013; Lim et al., 2015; Lim et al., 2014; Pedersen et al., 2020; Voronina et al., 2019).

These bacteriophages vary especially in terms of morphology, lytic activity and host range

but they all have the potential to control soft rot caused by *Pectobacterium* sp. as evidenced

by the semi-*in planta* experiments performed in these reports.

In this study, three lytic phages isolated from symptomatic tissues and environmental

samples, were characterized based on virion morphology, growth characteristics, and

physico-chemical properties. The lytic activity and efficacy of the phages to prevent tissue

maceration caused by *P. carotovorum* subsp. *carotovorum* in potato slices using monophage

and cocktail suspensions were also determined. The effect of the timing of phage application

on the efficacy of monophage treatments on whole potato tubers was also qualitatively

assessed. Based on the *in vitro* and semi-*in planta* assays, we report the biocontrol potential

of these lytic phages against bacterial soft rot caused by *P. carotovorum* subsp. *carotovorum.*

2. Materials and Methods

2.1.Bacterial Strains and Media

Pectobacterium carotovorum subsp. *carotovorum* BIOTECH 1752, obtained from the

Philippine National Collection of Microorganisms (National Institute of Molecular Biology

and Biotechnology, UPLB) was used as the primary host for phage isolation. A collection of

48 *Pectobacterium* spp. isolated from vegetable growing areas and market centers in the

Philippines (Host-Pathogen Interaction Laboratory, IWEP, UPLB), and eight other

representative strains from other genera (Microbiology Division, IBS, UPLB) were used for

the host range assay (Supplementary Table 1). For routine tests and maintenance, bacteria

were grown in Tryptone Soya Agar (TSA, HiMedia®, Mumbai, India) for 16-18 h at 30°C.

For liquid preparations, bacterial cultures were grown in Tryptone Soya Broth (TSB,

HiMedia®, Mumbai, India) for 12-18 h at 30°C with 125 rpm agitation.

2.2. Bacteriophage Isolation and Characterization

2.2.1. Isolation and Purification of Bacteriophages

Bacteriophages were isolated from rhizosphere soil, symptomatic tissues, and sewage

water (Czajkowski et al., 2014). Twenty grams (or mL) of each sample was

homogenized in 20 mL saline-magnesium (SM) buffer ($MgSO_4$, NaCl, $CaCl_2$, and

Tris-HCl, pH 7.5) and incubated at 30°C for 12-16 h. Five hundred μL of log phase (5

to 6-h subculture prepared from an overnight culture) *P. carotovorum* subsp.

carotovorum culture containing approximately 10^8 cells mL^{-1} (0.5 McFarland

Standard) was added to each enrichment set-up and was incubated as above.

Enrichment cultures were centrifuged at 3,000 rpm for 15 min and the supernatants

were chloroform-treated in 1:20 chloroform/supernatant ratio for at least 10 min. The

resulting lysates were screened for the presence of bacteriophages by spot test assay

(Mazzocco et al., 2009). Overlay mixtures, containing 4.5 mL soft agar (TSA), 0.5

mL 20 mM $CaCl_2$, and 300 μL of log phase *P. carotovorum* subsp. *carotovorum*

culture (0.5 MFS) were poured on Petri plates containing pre-solidified TSA. Five μL

of the phage lysates were spotted on the overlay and the spots were absorbed into the

agar for 10-15 min prior to incubation of plates at 30°C for 12-16 h. Zones of lysis on

the bacterial lawn indicated the presence of phages.

Phage purification was done using the double layer plaque assay (Kropinski et al.,

2009). Ten-fold dilutions were prepared from each positive lysate. In separate sterile

1.5-mL microcentrifuge tubes, 300 μL of diluted lysate suspensions were mixed with

300 μL log phase culture of *P. carotovorum* subsp. *carotovorum* (0.5 MFS) in TSB.

These were pre-incubated for 10 min at 4°C to allow adsorption of the phages. Molten

overlay (4.5 mL) mixed with 0.5 mL 20 mM $CaCl_2$ was added to the suspensions and

the resulting mixtures were poured onto Petri plates containing pre-solidified TSA.

The overlays were allowed to solidify for 15-20 min and the plates were incubated at

30°C for 12-16 h. Well-isolated plaques were picked using sterile pipette tips and the

phages were suspended in 500 µL SM buffer. Bacteriophage isolates were purified by

successive single plaque isolation for at least five times or until uniform plaque

morphology was observed.

2.2.2. *Plaque Formation Using Double Layer Plaque Assay*

Plaque morphologies were characterized after a series of double layer plaque assays

as described above (Jurczak-Kurek et al., 2016). The transparency and morphology of

the plaques formed by each bacteriophage isolate was described and plaque diameter

was measured using a digital caliper.

2.2.3. *Bacteriophage Lysate Preparation and Titration*

Double layer plaque assay plates with uniform plaques were flooded with 3-5 mL of

SM buffer to obtain the bacteriophage lysate. Plates were gently shaken for 15-20 min

and the overlay-SM buffer mixtures were collected and centrifuged at 8,000 rpm for

10 min. The supernatants were chloroform treated in 1:20 chloroform/lysate ratio and

were maintained at 4°C until titration. Lysates were subjected to double layer plaque

assay (or spot test) for titration as described above. Phage titer was determined and

expressed as plaque forming units (PFU mL^{-1}).

2.2.4. *Determination of Virion Morphology Using Transmission Electron Microscopy*

(TEM)

High titer lysates (at least 10^{10} PFU mL^{-1}) of the three phage isolates were submitted

to the Electron Microscopy Laboratory of the Research Institute for Tropical

Medicine (Muntinlupa City, Philippines). The phage suspension was placed on

carbon-coated copper grids, and 2% aqueous uranyl acetate was added for three min

to negatively stain the virions. Phages were visualized by transmission electron

microscopy (TEM; JEOLJEM-1220 Electron Microscope), and the images were

scanned with 800 ms exposure, 1.4 gain, and 1.0 bin with no sharpening, normal

contrast, and 1.00 gamma.

2.2.5. Optimal Multiplicity of Infection (MOI)

The optimal ratio of phage particles to the number of bacterial cells, also called the

optimal multiplicity of infection (MOI) was determined for the three phage isolates.

P. carotovorum subsp. *carotovorum* was infected with each phage isolate at four

different PFU mL^{-1}/cell mL^{-1} ratios (MOI): 0.01, 0.1, 1.0, and 10. Following

overnight incubation at 30°C, the suspensions were chloroform-treated (1:20) for 10

min and centrifuged at 8,000 rpm for 10 min. The supernatants were subjected to

double layer plaque assay to determine phage titer. The MOI demonstrating the

highest phage titer or PFU mL^{-1} was considered optimal.

2.2.6. Latent Period and Phage Burst Size

One-step growth experiments were conducted to determine phage latent period and

burst size following previously described protocol (Czajkowski et al., 2014). Cells of

P. carotovorum subsp. *carotovorum* at log phase were harvested by centrifugation at

12,000 rpm for 15 min at 10°C, and were resuspended in TSB to an OD$_{600}$ of 0.5 (c. 5

$\times$ 10^8 CFU mL^{-1}). Subsequently, 100 µL of 20mM CaCl$_2$ and 100 µL of the phage

isolate were added into the bacterial culture at MOI of 0.1 (10^7 PFU mL^{-1}/ 10^8 CFU

mL^{-1}). The phages were allowed to adsorb by incubation at 4°C for 10 min and the

suspensions were further incubated at 30°C for 10 min. The suspensions were

centrifuged at 12,000 rpm for 15 min at 10°C to remove unadsorbed phages. The

resulting pellet was resuspended in 10 mL fresh TSB. The mixtures were incubated at

30°C for 60 min with agitation at 125 rpm. Aliquots were taken from the mixtures

every 10 min and the titer was determined using the Spot Test method. The burst size

was calculated using the formula: $burst\ size = \frac{PFU_{final} - PFU_{initial}}{PFU_{initial}}$ and the

experiment was repeated twice with the same set-up and the results were averaged.

2.2.7. Adsorption Rate

Phage adsorption rate was determined following previously described protocol

(Czajkowski et al., 2014). One mL of *P. carotovorum* subsp. *carotovorum* (ca. 10^8

CFU mL^{-1}) was infected with each phage isolate by adding 100 μL of phage lysate

(ca. 10^8 pfu mL^{-1}) to reach MOI 0.1 and the resulting suspensions were incubated at

30°C for 20 min. After 0 (control), 1, 2, 4, 10 and 20 min, 50-μL aliquots were taken

from each suspension and added to 450 μL TSB. Ten-fold dilutions in SM buffer

were prepared and the residual titer of the unadsorbed phages were determined using

the spot test method as described above. The experiment was repeated twice with the

same set-up and the results were averaged. Phage adsorption was calculated as

follows: $\%adsorption = \frac{control\ titer - residual\ titer}{residual\ titer} x 100$.

2.2.8. Thermal Resistance and pH Stability

Thermal resistance of the three phages was determined at 30, 40, 50 and 60 °C.

Phages (10^8 PFU mL^{-1}) in SM buffer were incubated for 1 h at each temperature

condition and the surviving PFU was determined using double layer plaque assay

(Czajkowski et al., 2014).

The effect of pH on phage stability was determined using phages (10^8 PFU mL^{-1}) in

SM buffer within a pH range of 3.0 to 12.0. Phage lysates (10^8 PFU mL^{-1}) were added

to SM buffer, adjusted with 2N HCl or 3N NaOH to pH 3, 7, 9, and 12 (Lim et al.,

2013). The experiment was carried out by incubating the phage suspensions at 4°C for

16-24 h and the surviving PFU was determined using double layer plaque assay.

2.2.9.   Host Range Determination

The host ranges of the three phage isolates was determined using a collection of 48

Pectobacterium spp. isolates and eight representative strains from different genera

(Supplementary Table 1) using the spot test assay. *Pectobacterium* spp. were isolated

in the years 2017-2019 from symptomatic tissues of various vegetable crops.

For the spot test assay, bacterial cultures were grown in TSB at 30°C for 12-16 h with

agitation at 125 rpm. From the resulting bacterial cultures, 300 μL (0.5 MFS) of each

was inoculated into the overlay agar and was subsequently poured into a prepared

TSA plate and allowed to solidify at room temperature for 30 min. Five μL of the

phage lysate (10^9 PFU mL^{-1}) was spotted onto the top agar layer, and the plates were

allowed to dry at room temperature for 10-15 min. Following overnight incubation at

30°C, plates were inspected for lysis zones.

2.3. In Vitro Challenge Test

Bacterial inactivation using monophage (vB_PcaP-A3, vB_PcaM-D1, or vB_PcaM-J3)

and cocktail treatments (vB_PcaP-A3+D1, vB_PcaP-A3+J3,   vB_PcaM-D1+J3, or

vB_PcaP-A3+D1+J3) was determined following the method described by Chen et al.

(2018). For monophage treatments, log phase culture of *P. carotovorum* subsp.

carotovorum (OD_{600} of 0.5, c. 5×10^8 CFU mL^{-1}) was mixed with the phage lysate at

optimal MOI. For cocktail preparations, low (0.001) and high (100) MOI were used for

the experiment. The bacteria-phage mixtures were incubated at 30°C with agitation at 125

rpm and the OD readings at 600nm were recorded (SPECTROstar^{Nano}, Ortenberg,

Germany) at 15-minute intervals for eight hours to monitor the changes in bacterial

density.

2.4. Semi-In Planta Assays

2.4.1. Phage Biocontrol Using Monophage and Cocktail Treatments on Tissue Slices

The efficacies of monophage and cocktail treatments against soft rot caused by *P.*

carotovorum subsp. *carotovorum* were evaluated using a potato slice assay

(Czajkowski et al., 2015). Potato tubers (cultivar Igorota) were surface sterilized by

soaking in 70% ethanol for 10 min, washed with sterile distilled water, and air dried

under the laminar flow hood. The tubers were cut into 7.0 mm-thick transverse disks

using sterile knives and three wells (5-mm diameter) were made on each slice using a

sterile cork borer. For monophage treatments, log phase culture of *P. carotovorum*

subsp. carotovorum containing approximately 10^8 cells mL^{-1} (0.5 MFS) was mixed

with the phage lysates at optimal MOI. For cocktail preparations, low (0.001) and

high (100) MOIs were used for the experiment. Three potato slices were used for each

treatment. Wells were filled with 20 µL of the bacteria-phage mixtures, 20 µL of

sterile SM buffer as negative control, and 20 µL of *P. carotovorum* subsp.

carotovorum (0.5 MFS) as positive control. The diameter of the rotten tissues around

the wells inoculated with bacteria and wells co-inoculated with bacteria and phages

were measured after incubation at 30°C for 72 h. Treatment means were compared

using one-way ANOVA with Tukey's post-test ($\alpha = 0.05$).

 2.4.2.   Protective and Curative Potential of Monophage Treatments against Soft Rot

A potato tuber assay was conducted to determine the protective and curative effect of

monophage treatments at various application time points relative to *P. carotovorum*

subsp. *carotovorum* inoculation following a previously described protocol by

Czajkowski et. al. (2014). Potato tubers (cultivar Igorota) were washed in running

water to remove soil and other organic particles, surface-sterilized with 70% ethanol

for 10 min, rinsed with sterile water for three times, and air dried under laminar

flowhood. For each tuber, two wells (8 mm diameter) were made using flame-

sterilized cork borer. Two set-ups were prepared for the experiment: (1) advanced

phage introduction (API) treatment, and (2) delayed phage introduction (DPI)

treatment. For both treatments, log phase culture of *P. carotovorum subsp.*

carotovorum containing approximately 10^7 cells mL^{-1} was used and the phage lysate

was added to the wells at MOI 10. In the API treatment, wells were filled with 25 μL

of the phage suspension at 0, 1, 3, 8, 15 and 30 h prior to the inoculation of 25 μL of

the bacterial suspension. In the DPI treatment, wells were filled with 25 μL of the

phage suspension at 0, 1, 3, 8, 15 and 30 h after bacterial inoculation. Three potato

tubers were used per time point for each of the API and DPI treatment. For the control

set-ups, 25 μL aliquots of sterile water was used as negative control while 25 μL of *P.*

carotovorum subsp. *carotovorum* suspension was used as positive control. The tubers

were placed in a humid box and the degree of tissue maceration was qualitatively

assessed (scored as +/-) after incubation at 30°C for 72 h.

3. Results

3.1.Isolation and Characterization of Bacteriophages

3.1.1. Isolation of Lytic Bacteriophages

Bacteriophages vB_PcaP-A3, vB_PcaM-D1 and vB_PcaM-J3 were isolated from

rhizosphere soil, symptomatic tissues, and sewage water, respectively, using *P.*

carotovorum subsp. *carotovorum* as enrichment and isolation host. The three phages

were named following the unified phage naming system proposed by Kropinski et al.

(2009). All three phages formed clear plaques typical for lytic phages, with varying

sizes (Table 1).

3.1.2. Virion Morphology

Examination of the phages using TEM revealed that vB_PcaP-A3 exhibited

podovirus morphology while vB_PcaM-D1 and vB_PcaM-J3 showed virion

morphology similar to members of the myovirus group (Table 1, Figure 1). Phage

vB_PcaP-A3 has an icosahedral head of 80 nm and short non-contractile tail of 20 nm

(Table 1). Phages vB_PcaM-D1 and vB_PcaM-J3 both have icosahedral heads of

around 75 nm, with longer, rigid contractile tails that are about 95-135 nm (Table 1,

Figure 1).

3.1.3. Characterization of Bacteriophages

All three phages exhibited strong adsorption to *P. carotovorum* subsp. *carotovorum,*

and approximately 90% of the phage particles adsorbed to the bacterial cells within

five min (Figure 2). vB_PcaP-A3 and vB_PcaM-D1 showed similar infection kinetics

with a latent period of 10 min and a burst size of 142-151 progeny phages/cell,

whereas vB_PcaM-J3 showed a longer latent period of 20 min and a smaller burst

size of 42 progeny phages/cell (Table 1, Figure 2A & 2B). Among the three phages,

vB_PcaP-A3 has the lowest optimal MOI (0.01), followed by vB_PcaM-D1 (1.0) and

vB_PcaM-J3 (10.0) (Table 1). All of the three phages retained infectivity after

incubation at 30°C, 40°C, and 50°C for 1 h, and in the pH range 3-9 for 16-24 h

(Figure 2C) *[Figure 1 near here; Table 1 in the next page].*

[Figure 2 near here]

3.1.4. Host Range

Spot test assay revealed that the three phages showed antibacterial activity specific for

P. carotovorum subsp. *carotovorum.* None of the three phages was able to lyse the 48

Pectobacterium spp. isolates and the eight representative strains from other genera.

3.1.5. Phage Activity In Vitro

Killing curves established for vB_PcaP-A3, vB_PcaM-D1 and vB_PcaM-J3 showed

that all three phages have strong lytic activity and were highly effective against *P.*

carotovorum subsp. *carotovorum* at optimal MOI (Figure 3A). The bacterial cell

density remained less than 0.05 (OD_{600}) throughout the 8-h observation period. The

absence of any increase in OD_{600} reading also suggests that there is no observed phage

resistance development of the host, at least during the observation period (Muturi et

al., 2019). These results indicate that monophage treatments of vB_PcaP-A3,

vB_PcaM-D1 and vB_PcaM-J3 at optimal MOI were equally effective in inhibiting

the growth of *P. carotovorum* subsp. *carotovorum in vitro*.

For the cocktail treatments, low MOI (0.001) and high MOI (100) were used in the *in*

vitro challenge tests. Each phage was used at the same concentration and the phage

cocktails consisting of two or three phages were combined and evaluated. At MOI of

0.001, the killing curves of the phage cocktails composed of two or three phages

maintained its bacterial inactivation with an $OD_{600} < 0.05$ for 8 h (Figure 4A). At low

MOI, it appears that a phage cocktail containing any of the three phages effectively

controlled bacterial growth during the 8-h observation period. At MOI 100, killing

curves showed that all cocktail treatments were highly effective against *P.*

carotovorum subsp. *carotovorum* (Figure 4A). The bacterial density remained less

than 0.05 (OD_{600}) throughout the duration of the experiment.

3.1.6.   Phage-Mediated Biocontrol Using Monophage and Cocktail Treatments

3.1.6.1. Monophage and cocktail treatments reduced soft rot symptoms in tissue slices

The average diameter of rotting tissue was significantly lower in all monophage

treatments than in the set-up inoculated with *P. carotovorum* subsp. *carotovorum* only

(Figure 3B-D). The set-up co-inoculated with *P. carotovorum* subsp. *carotovorum*

and vB_PcaP-A3 exhibited the lowest diameter of rotting tissue among the three

treatments (Figure 3B-D). Phage vB_PcaP-A3 was able to reduce potato tuber tissue

maceration by at least 60%, while phages vB_PcaM-D1 and vB_PcaM-J3 were able

to reduce tissue maceration by at least 50% of that observed for the control potato

slices inoculated with *P. carotovorum* subsp. *carotovorum* only (Figure 3B-D). All of

the phage cocktail treatments completely prevented tissue maceration at low and high

MOI (Figure 4B-D).

3.1.6.2. Advanced phage introduction can protect tubers from soft rot

The potential use of phages as a post-harvest disease control strategy was assessed

using potato tubers artificially inoculated with *P. carotovorum* subsp. *carotovorum*

treated with monophage suspensions applied at different time points. Results of the

whole tuber assay showed that in the API treatment, monophage suspensions of the

three phages applied on to the wound up to 30 h before bacterial inoculation provided

protection against soft rot disease development (Figure 3E). In the DPI treatment, a 3-

h and 8-h delay in the application of monophage suspensions of vB_PcaM-J3, and

vB_PcaP-A3 and vB_PcaM-D1, respectively, following bacterial inoculation resulted

in disease development (Figure 3E). *[Figure 3 near here]*

[Figure 4 near here]

4. Discussion

Phage-mediated biocontrol of soft rot disease is a promising and an up-and-coming

technology that has been widely explored worldwide. To our knowledge, this is the first

report on the isolation and characterization of bacteriophages infecting *P. carotovorum* subsp.

carotovorum in the Philippines, which explores some aspects that need to be considered for

potential formulation of a postharvest biocontrol product against soft rot disease including

phage stability, bioactivity in tissues, and application timing.

Three lytic bacteriophages infecting *P. carotovorum* subsp. *carotovorum* designated as

vB_PcaP-A3, vB_PcaM-D1, and vB_PcaM-J3, were isolated from environmental samples

including rhizosphere soil, symptomatic tissues, and sewage water. TEM analysis of the three

phages revealed that vB_PcaP-A3 has a podovirus morphology similar to *Pectobacterium*

phage Arno160, *Pectobacterium* phage vB_PatP_CB4, and *Dickeya* phage Amaethon

(Buttimer, Hendrix, et al., 2018; Shneider et al., 2020). Meanwhile, phages vB_PcaM-D1 and

vB_PcaM-J3 exhibited myoviral morphology similar to *Ackermannviridae Dickeya* phage

PP35, *Pectobacterium* phage PP101, and *Pectobacterium* phage vB_PcaM_CBB (Kabanova

et al., 2019; Lukianova et al., 2020).

The latent period and burst size of phages vB_PcaP-A3, vB_PcaM-D1, and vB_PcaM-J3

were determined based on one-step growth curve experiments. The phages were further

characterized, and the adsorption rate, optimal MOI, and stability on varying temperature and

pH levels were also determined. Of the three phage isolates, vB_PcaP-A3 has the lowest

optimal MOI of 0.01, a latent period of 10 min and a calculated burst size value of 142

PFU/cell. Phage vB_PcaM-D1 has an optimal MOI of 0.1, latent period of 10 min and a burst

size of 151 PFU/cell. vB_PcaM-J3 has an optimal MOI of 10, longest latent period of 30 min,

and the lowest calculated burst size of 42 PFU/cell. Since phages displaying shorter latent

periods and higher burst sizes are reported to provide better efficiency than phages with

longer latent periods in cultures where the density of the host is high, vB_PcaM-J3 showed

less promising growth characteristics compared to vB_PcaP-A3 and vB_PcaM-D1 (Abedon

et al., 2003). All three phages, however, showed strong adsorption to their host evidenced by

the observed 90% phage adsorption within 5 min after incubation. When their stabilities on

varying temperature and pH levels were tested, all three phages survived incubations at 30°C-

50°C and at pH 3-9, but were inactivated at 60°C and at pH 12. These are important

environmental factors in biological control applications. Since temperatures under field

conditions do not exceed 50°C and pH levels usually range from pH 3.3-4.3, results indicate

that vB_PcaP-A3, vB_PcaM-D1 and vB_PcaM-J3 can be suitable for possible field

applications (Navarrete et al., 2017).

All three phages showed specificity to their isolation host, *P. carotovorum* subsp.

carotovorum. Phages are generally known to have narrow host ranges, typically limited to

strains within a certain species of their bacterial host (Ross et al., 2016). This can allow the

formulation of phage cocktails targeting bacterial species of interest only, which may be a

certain bacterial plant pathogen or a specific bacterium in a microbial community that

negatively impacts plant growth or health conditions. Developing a phage-based biocontrol

product that can eliminate multiple genera, or even different related species could be

challenging, but combining phages specific to target hosts may overcome this problem

(Buttimer et al., 2017; Frampton et al., 2012). The specificity of phages also reduces the risk

of harming the natural microbiota, especially those non-target bacteria that are potentially

beneficial, and it limits the emergence of phage-resistant bacteria (Loc-Carrillo & Abedon,

2011).

In recent years, there were numerous attempts to systematize the single and cocktail

application of phages to tubers and vegetable crops. For example, the first bacteriophage of

P. carotovorum subsp. *carotovorum* that was sequenced, PP1, exhibited effective protection

against its host in lettuce (Lim et al., 2013). In another study, two other phages namely

□PD10.3 and □PD23.1 were applied together in both whole potato tubers and slices against

a number of *Pectobacterium* and *Dickeya* isolates. Tissue maceration was significantly

reduced by 80% and 95% in slices and whole tubers, respectively (Czajkowski et al., 2015).

Two further studies revealed effective inhibition of SRPs in potato tubers using monophage

formulations (J. Lee et al., 2017; Smolarska et al., 2018). In this study, the efficacy of the

three bacteriophages, vB_PcaP-A3, vB_PcaM-D1, and vB_PcaM-J3, in both monophage and

cocktail suspensions (combinations of two and three phages) were assessed through *in vitro*

and semi-*in plant*a bacterial challenge tests, and we found that both monophage and phage

cocktails completely inactivated the growth of *P. carotovorum* subsp. *carotovorum.*

However, phages in cocktail suspensions were more effective over individual application in

terms of their protective effect against soft rot by *P. carotovorum* subsp. *carotovorum* as

indicated by the complete maceration reduction in both MOI (0.001 and 100) tested in the

study. Recent reports of phage cocktail formulations against *P. carotovorum* subsp.

carotovorum have also described efficient inhibition of bacterial growth both *in vitro* and *in*

vivo (Carstens et al., 2019; Muturi et al., 2019; Zaczek-Moczydłowska et al., 2020). The use

of higher MOI in phage formulations has also been reported to increase the chance for the

bacteriophages to infect all host cells, thus effectively killing all host cells after one round of

replication (Alves et al., 2016; Thomas, 2001).

Using different novel phage isolates against a number of SRP model strains, our work and the

several studies mentioned above have provided preliminary evidence of the biocontrol

potential of bacteriophages in controlling bacterial soft rot. However, a comprehensive

understanding of their bioefficacy in postharvest applications is still limited and our work

contributes to this pool of information by determining the efficient timing of phage treatment

on tissues.

Preliminary experiments conducted to assess possible timing strategies in employing phage

treatments to control post-harvest soft rot revealed the protective effect of introducing phages

prior to bacterial inoculation (Figure 3E). In each monophage treatment, advanced phage

introduction (API) treatments of up to 30 h were observed to protect the tubers from

developing the disease (Figure 3E). With API, we hypothesize that, considering that the

phage number applied is higher than or equal to the optimal MOI, the phages were able to

rapidly kill the inoculated bacterial population, thus suppressing both the bacterial growth

and the emergence of visible symptoms of the disease.

In contrast, a shorter window of time wherein the phage treatment seemed to offer effective

protection from the disease was observed in the delayed phage introduction (DPI) treatment.

This observation, in line with the hypothesized phage-bacterium interaction in the API

treatment, seemed to suggest that a critical bacterial population relative to the phage

population was required to effect noticeable disease manifestation. It was previously reported

that quorum sensing regulates the coordination of the synthesis of virulence factors, including

the production of plant cell wall degrading enzymes by *Pectobacterium* spp, which links the

observed longer protective window of API treatment and the reduced protective window

observed in DPI treatment (Pollumaa et al., 2012).

Furthermore, taking both into account the differences in the characteristics of the phages and

the MOI used in this experiment (MOI 10.0, relative to bacterial population at time 0), it

follows that, for the DPI treatment, a shorter protective window shall be observed for

vB_PcaM-J3, which has the longest latent period (30 min), lowest burst size (42 CFU/cell)

and highest MOI (10.0). That is, with the advanced introduction of the bacterial population of

at least 3 h, the bacterial population seemed to have grown enough such that the vB_PcaM-J3

population is not at its optimal MOI anymore. In line with this is the observed longer

protective window afforded by vB_PcaP-A3 and vB_PcaM-D1 that have lower optimal

MOIs of 0.01 and 1.0 respectively, which are both lower than the MOI used in this

experiment, and higher burst sizes of 151 CFU/cell and 142 CFU/cell, respectively. These

characteristics may have also provided a buffering effect to account for the bacterial growth

in DPI treatment.

Taken together, these results are suggestive of a potential post-harvest soft rot disease control

strategy that may maximize the benefit of phage treatment by introducing efficiently lytic

phages (ie. short latent period, larger burst size, and low optimal MOI) at a high MOI at the

earliest time point after harvest. By doing so, while a small population of the soft rot-causing

bacteria may have been inadvertently introduced during the process of harvesting, this

population can be effectively controlled thus preventing bacterial growth, disease
manifestation, and cross-contamination.

However, we are also aware that the use of higher MOI could also possibly increase the
emergence of phage-resistant bacterial variants (Chen et al., 2018). This is due to the rapid
evolution of the bacterial population from the dominance of phage-sensitive clones to phage-
resistant clones when exposed to the selection pressure of lytic phages (Castillo et al., 2015;
Hyman & Abedon, 2010; Middelboe, 2000; Middelboe et al., 2001). This implies that the use
of heavy phage loads is an important selection factor that should be taken into account in
phage therapy and that lower MOIs should also be used when testing protective effect and
bacterial inactivation by phages, particularly in actual plant tissues (Chen et al., 2018). To
this end, further experimental studies to optimize the phage treatment parameters and to
identify the minimum effective phage number for potential field and post-harvest application
are to be conducted.

Data statement

All datasets presented in this paper are included in the article.

CRediT authorship contribution statement

Aeron Jade S. Parena: Methodology, Validation, Formal Analysis, Investigation, Data
Curation, Writing-Original Draft, Writing-Review & Editing, Visualization. **Benji Brayan I.**
Silva: Conceptualization, Methodology, Validation, Formal Analysis, Writing-Review &
Editing. **Rae Mark L. Mercado:** Methodology, Validation, Investigation, Data Curation.
Adrian Adelbert A. Sendon: Methodology, Validation, Investigation. **Freddiewebb B.**
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Supervision, Project Administration, Funding Acquisition. **Jaymee R. Encabo:**

Conceptualization, Data Curation, Writing-Review & Editing, Supervision, Project

administration, Funding acquisition.

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Declaration of competing interest

491　The authors declare that the research was conducted with no known competing financial

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References

Abedon, S. T., Hyman, P., & Thomas, C. (2003). Experimental examination of bacteriophage latent-period evolution as a response to bacterial availability. *Applied and Environmental Microbiology*, *69*(12), 7499-7506. https://doi.org/10.1128/AEM.69.12.7499-7506.2003

Adeolu, M., Alnajar, S., Naushad, S., & Gupta, R. S. (2016). Genome-based phylogeny and taxonomy of the 'Enterobacteriales': proposal for *Enterobacterales* ord. nov. divided

into the families *Enterobacteriaceae*, *Erwiniaceae* fam. nov., *Pectobacteriaceae* fam.
nov., *Yersiniaceae* fam. nov., *Hafniaceae* fam. nov., *Morganellaceae* fam. nov., and
Budviciaceae fam. nov. *International Journal of Systematic and Evolutionary*
Microbiology, *66*(12), 5575-5599. https://doi.org/10.1099/ijsem.0.001485
Agyemang, P. A., Kabir, M. N., Kersey, C. M., & Dumenyo, C. K. (2020). The bacterial soft
rot pathogens, *Pectobacterium carotovorum* and *P. atrosepticum*, respond to different
classes of virulence-inducing host chemical signals. *Horticulturae*, *6*(1), 1-13.
https://doi.org/10.3390/horticulturae6010013
Alves, D. R., Perez-Esteban, P., Kot, W., Bean, J. E., Arnot, T., Hansen, L. H., Enright, M.
C., & Jenkins, A. T. A. (2016). A novel bacteriophage cocktail reduces and disperses
Pseudomonas aeruginosa biofilms under static and flow conditions. *Microbial*
Biotechnology, *9*(1), 61-74. https://doi.org/10.1111/1751-7915.12316
Bayogan, E. R. V., Jimenez, E. F., Boteng, J. D., Bautista, O. K., & Macario, C. B. (2011).
Improving postharvest cabbage (*Brassica oleracea* L. var. *capitata*) quality using
alum   and   newspaper   wrap.   *Banwa*,   *6*(2),   1-11.
https://doi.org/10.3860/banwa.v6i2.1964
Behlau, F., Canteros, B. I., Minsavage, G. V., Jones, J. B., & Graham, J. H. (2011).
Molecular characterization of copper resistance genes from *Xanthomonas citri* subsp.
citri and *Xanthomonas alfalfae* subsp. *citrumelonis*. *Applied and Environmental*
Microbiology, *77*(12), 4089-4096. https://doi.org/10.1128/AEM.03043-10
Bhat, K. A., Masood, S. D., Bhat, N. A., Bhat, M. A., Razvi, S. M., Mir, M. R., Akhtar, S.,
Wani, N., & Habib, M. (2010). Current status of post harvest soft rot in vegetables: a
review.   *Asian   Journal   of   Plant   Sciences*,   *9*(4),   200-208.
https://doi.org/10.3923/ajps.2010.200.208
Buttimer, C., Hendrix, H., Lucid, A., Neve, H., Noben, J. P., Franz, C., O'Mahony, J.,
Lavigne, R., & Coffey, A. (2018). Novel N4-like bacteriophages of *Pectobacterium*
atrosepticum.   *Pharmaceuticals*   *(Basel)*,   *11*(2),   45.
https://doi.org/10.3390/ph11020045
Buttimer, C., Lucid, A., Neve, H., Franz, C., O'Mahony, J., Turner, D., Lavigne, R., &
Coffey, A. (2018). *Pectobacterium atrosepticum* phage vB_PatP_CB5: a member of
the   proposed   genus   '*Phimunavirus*'.   *Viruses*,   *10*(8),   394.
https://doi.org/10.3390/v10080394
Buttimer, C., Lynch, C., Hendrix, H., Neve, H., Noben, J. P., Lavigne, R., & Coffey, A.
(2020). Isolation and characterization of *Pectobacterium* phage vB_PatM_CB7: new
insights   into   the   genus   *Certrevirus*.   *Antibiotics   (Basel)*,   *9*(6),   1-24.
https://doi.org/10.3390/antibiotics9060352
Buttimer, C., McAuliffe, O., Ross, R. P., Hill, C., O'Mahony, J., & Coffey, A. (2017).
Bacteriophages and bacterial plant diseases. *Frontiers in Microbiology*, *8*, 34.
https://doi.org/10.3389/fmicb.2017.00034
Carstens, A. B., Djurhuus, A. M., Kot, W., & Hansen, L. H. (2019). A novel six-phage
cocktail reduces *Pectobacterium atrosepticum* soft rot infection in potato tubers under
simulated   storage   conditions.   *FEMS   Microbiology   Letters*,   *366*(9),   1-8.
https://doi.org/10.1093/femsle/fnz101
Castillo, D., Christiansen, R. H., Dalsgaard, I., Madsen, L., & Middelboe, M. (2015).
Bacteriophage   resistance   mechanisms   in   the   fish   pathogen   *Flavobacterium*
psychrophilum: linking genomic mutations to changes in bacterial virulence factors.
Applied   and   Environmental   Microbiology,   *81*(3),   1157-1167.
https://doi.org/10.1128/AEM.03699-14

Charkowski, A. (2018). The changing face of bacterial soft-rot diseases. *Annual Review of Phytopathology*, *56*(1), 269-288. https://doi.org/10.1146/annurev-phyto-080417-045906

Charkowski, A., Blanco, C., Condemine, G., Expert, D., Franza, T., Hayes, C., Hugouvieux-Cotte-Pattat, N., Lopez Solanilla, E., Low, D., Moleleki, L., Pirhonen, M., Pitman, A., Perna, N., Reverchon, S., Rodriguez Palenzuela, P., San Francisco, M., Toth, I., Tsuyumu, S., van der Waals, J., van der Wolf, J., Van Gijsegem, F., Yang, C. H., & Yedidia, I. (2012). The role of secretion systems and small molecules in soft-rot *Enterobacteriaceae* pathogenicity. *Annual Review of Phytopathology*, *50*, 425-449. https://doi.org/10.1146/annurev-phyto-081211-173013

Chen, L., Yuan, S., Liu, Q., Mai, G., Yang, J., Deng, D., Zhang, B., Liu, C., & Ma, Y. (2018). In vitro design and evaluation of phage cocktails against *Aeromonas salmonicida*. *Frontiers in Microbiology*, *9*(JUL), 1-13. https://doi.org/10.3389/fmicb.2018.01476

Cooksey, D. A. (1990). Genetics of bactericide resistance in plant pathogenic bacteria. *Annual Review of Phytopathology*, *28*(1), 201-219. https://doi.org/10.1146/annurev.py.28.090190.001221

Czajkowski, R. (2015). Bacteriophages of soft rot *Enterobacteriaceae*-a minireview. *FEMS Microbiology Letters*, *363*(2), 1-6. https://doi.org/10.1093/femsle/fnv230

Czajkowski, R., Ozymko, Z., de Jager, V., Siwinska, J., Smolarska, A., Ossowicki, A., Narajczyk, M., & Lojkowska, E. (2015). Genomic, proteomic and morphological characterization of two novel broad host lytic bacteriophages PhiPD10.3 and PhiPD23.1 infecting pectinolytic *Pectobacterium* spp. and *Dickeya* spp. *PLOS ONE*, *10*(3), e0119812. https://doi.org/10.1371/journal.pone.0119812

Czajkowski, R., Ozymko, Z., & Lojkowska, E. (2014). Isolation and characterization of novel soilborne lytic bacteriophages infecting *Dickeya* spp. biovar 3 ('*D. solani*'). *Plant Pathology*, *63*(4), 758-772. https://doi.org/10.1111/ppa.12157

Czajkowski, R., Pérombelon, M. C. M., Van Veen, J. A., & Van der Wolf, J. M. (2011). Control of blackleg and tuber soft rot of potato caused by *Pectobacterium* and *Dickeya* species: a review. *Plant Pathology*, *60*(6), 999-1013. https://doi.org/10.1111/j.1365-3059.2011.02470.x

Davidsson, P. R., Kariola, T., Niemi, O., & Palva, E. T. (2013). Pathogenicity of and plant immunity to soft rot pectobacteria. *Frontiers in Plant Science*, *4*, 191. https://doi.org/10.3389/fpls.2013.00191

Frampton, R. A., Pitman, A. R., & Fineran, P. C. (2012). Advances in bacteriophage-mediated control of plant pathogens. *International Journal of Microbiology*, *2012*, 1-11, Article 326452. https://doi.org/10.1155/2012/326452

Gardan, L., Gouy, C., Christen, R., & Samson, R. (2003). Elevation of three subspecies of *Pectobacterium carotovorum* to species level: *Pectobacterium atrosepticum* sp. nov., *Pectobacterium betavasculorum* sp. nov. and *Pectobacterium wasabiae* sp. nov. *International Journal of Systematic and Evolutionary Microbiology*, *53*(2), 381-391. https://doi.org/10.1099/ijs.0.02423-0

Hadas, H., Einav, M., Fishov, I., & Zaritsky, A. (1997). Bacteriophage T4 development depends on the physiology of its host *Escherichia coli*. *Microbiology*, *143*(1), 179-185. https://doi.org/10.1099/00221287-143-1-179

Hyman, P., & Abedon, S. T. B. T. A. i. A. M. (2010). Chapter 7 - Bacteriophage host range and bacterial resistance ((Vol. 70, pp. 217-248). Academic Press. https://doi.org/https://doi.org/10.1016/S0065-2164(10)70007-1

Jones, J. B., Jackson, L. E., Balogh, B., Obradovic, A., Iriarte, F. B., & Momol, M. T. (2007). Bacteriophages for plant disease control. *Annual Review of Phytopathology*, *45*, 245-262. https://doi.org/10.1146/annurev.phyto.45.062806.094411

Jones, J. B., Vallad, G. E., Iriarte, F. B., Obradovic, A., Wernsing, M. H., Jackson, L. E., Balogh, B., Hong, J. C., & Momol, M. T. (2012). Considerations for using bacteriophages for plant disease control. *Bacteriophage*, *2*(4), 208-214. https://doi.org/10.4161/bact.23857

Jurczak-Kurek, A., Gasior, T., Nejman-Faleńczyk, B., Bloch, S., Dydecka, A., Topka, G., Necel, A., Jakubowska-Deredas, M., Narajczyk, M., Richert, M., Mieszkowska, A., Wróbel, B., Węgrzyn, G., & Węgrzyn, A. (2016). Biodiversity of bacteriophages: morphological and biological properties of a large group of phages isolated from urban sewage. *Scientific Reports*, *6*(1), 1-17. https://doi.org/10.1038/srep34338

Kabanova, A. P., Shneider, M. M., Korzhenkov, A. A., Bugaeva, E. N., Miroshnikov, K. K., Zdorovenko, E. L., Kulikov, E. E., Toschakov, S. V., Ignatov, A. N., Knirel, Y. A., & Miroshnikov, K. A. (2019). Host specificity of the *Dickeya* bacteriophage PP35 is directed by a tail spike interaction with bacterial O-antigen, enabling the infection of alternative non-pathogenic bacterial host. *Frontiers in Microbiology*, *9*, 1-11. https://doi.org/10.3389/fmicb.2018.03288

Kalischuk, M., Hachey, J., & Kawchuk, L. (2015). Complete genome sequence of phytopathogenic *Pectobacterium atrosepticum* bacteriophage Peat1. *Genome Announcements*, *3*(4), 1-2. https://doi.org/10.1128/genomeA.00760-15

Kropinski, A. M., Mazzocco, A., Waddell, T. E., Lingohr, E., & Johnson, R. P. (2009). Enumeration of bacteriophages by double agar overlay plaque assay. *Methods in molecular biology (Clifton, N.J.)*, *501*, 69-76. https://doi.org/10.1007/978-1-60327-164-6_7

Lee, D. H., Lee, J.-H., Shin, H., Ji, S., Roh, E., Jung, K., Ryu, S., Choi, J., & Heu, S. (2012). Complete genome sequence of *Pectobacterium carotovorum* subsp. *carotovorum* bacteriophage My1. *Journal of Virology*, *86*(20), 11410-11411. https://doi.org/10.1128/jvi.01987-12

Lee, J., Kim, S., & Park, T. H. (2017). Diversity of bacteriophages infecting *Pectobacterium* from potato fields. *Journal of Plant Pathology*, *99*(2), 453-460. http://www.jstor.org/stable/44686791

Li, X., Fu, L., Chen, C., Sun, W., Tian, Y., & Xie, H. (2019). Characteristics and rapid diagnosis of *Pectobacterium carotovorum* spp. associated with bacterial soft rot of vegetables in China. *Plant Disease*, *104*(4), 1158-1166. https://doi.org/10.1094/PDIS-05-19-1033-RE

Liao, C.-H., McEvoy, J. L., & Smith, J. L. (2003). Control of bacterial soft rot and foodborne human pathogens on fresh fruits and vegetables. *Advances in plant Disease Management*, *661*(2), 165-193.

Lim, J. A., Heu, S., Park, J., & Roh, E. (2017). Genomic characterization of bacteriophage vB_PcaP_PP2 infecting *Pectobacterium carotovorum* subsp. *carotovorum,* a new member of a proposed genus in the subfamily *Autographivirinae*. *Archives of Virology*, *162*(8), 2441-2444. https://doi.org/10.1007/s00705-017-3349-6

Lim, J. A., Jee, S., Lee, D. H., Roh, E., Jung, K., Oh, C., & Heu, S. (2013). Biocontrol of *Pectobacterium carotovorum* subsp. *carotovorum* using bacteriophage PP1. *Journal of Microbiology and Biotechnology*, *23*(8), 1147-1153. https://doi.org/10.4014/jmb.1304.04001

Lim, J. A., Lee, D. H., & Heu, S. (2015). Isolation and genomic characterization of the T4-like bacteriophage PM2 infecting *Pectobacterium carotovorum* subsp. *carotovorum*. *Plant Pathology Journal*, *31*(1), 83-89. https://doi.org/10.5423/PPJ.NT.09.2014.0099

Lim, J. A., Shin, H., Lee, D. H., Han, S. W., Lee, J. H., Ryu, S., & Heu, S. (2014). Complete genome sequence of the *Pectobacterium carotovorum* subsp. *carotovorum* virulent bacteriophage PM1. *Archives of Virology*, *159*(8), 2185-2187. https://doi.org/10.1007/s00705-014-2005-7

Loc-Carrillo, C., & Abedon, S. T. (2011). Pros and cons of phage therapy. *Bacteriophage*, *1*(2), 111-114. https://doi.org/10.4161/bact.1.2.14590

Lukianova, A. A., Shneider, M. M., Evseev, P. V., Shpirt, A. M., Bugaeva, E. N., Kabanova, A. P., Obraztsova, E. A., Miroshnikov, K. K., Senchenkova, S. N., Shashkov, A. S., Toschakov, S. V., Knirel, Y. A., Ignatov, A. N., & Miroshnikov, K. A. (2020). Morphologically different *Pectobacterium brasiliense* bacteriophages PP99 and PP101: deacetylation of O-polysaccharide by the tail spike protein of phage PP99 accompanies the infection. *Frontiers in Microbiology*, *10*, 1-15. https://doi.org/10.3389/fmicb.2019.03147

Mattinen, L., Nissinen, R., Riipi, T., Kalkkinen, N., & Pirhonen, M. (2007). Host-extract induced changes in the secretome of the plant pathogenic bacterium *Pectobacterium atrosepticum*. *Proteomics*, *7*(19), 3527-3537. https://doi.org/10.1002/pmic.200600759

Mazzocco, A., Waddell, T. E., Lingohr, E., & Johnson, R. P. (2009). Enumeration of bacteriophages using the small drop plaque assay system. *Methods in molecular biology (Clifton, N.J.)*, *501*, 81-85. https://doi.org/10.1007/978-1-60327-164-6_9

McManus, P. S., Stockwell, V. O., Sundin, G. W., & Jones, A. L. (2002). Antibiotic use in plant agriculture. *Annual Review of Phytopathology*, *40*, 443-465. https://doi.org/10.1146/annurev.phyto.40.120301.093927

Middelboe, M. (2000). Bacterial growth rate and marine virus–host dynamics. *Microbial Ecology*, *40*(2), 114-124. https://doi.org/10.1007/s002480000050

Middelboe, M., Hagström, A., Blackburn, N., Sinn, B., Fischer, U., Borch, N. H., Pinhassi, J., Simu, K., & Lorenz, M. G. (2001). Effects of bacteriophages on the population dynamics of four strains of pelagic marine bacteria. *Microbial Ecology*, *42*(3), 395-406. https://doi.org/10.1007/s00248-001-0012-1

Monjil, M. S., Hasan, M. M., & Hoque, M. A. (2021). Estimation of post-harvest losses of potato in markets of some selected districts in Bangladesh. *Food and Agri Economics Review*, *1*(1), 21-27. https://doi.org/10.26480/faer.01.2021.21.27

Muturi, P., Yu, J., Maina, A. N., Kariuki, S., Mwaura, F. B., & Wei, H. (2019). Bacteriophages isolated in China for the control of *Pectobacterium carotovorum* causing potato soft rot in Kenya. *Virologica Sinica*, *34*(3), 287-294. https://doi.org/10.1007/s12250-019-00091-7

Nabhan, S., De Boer, S. H., Maiss, E., & Wydra, K. (2013). *Pectobacterium aroidearum* sp. nov., a soft rot pathogen with preference for monocotyledonous plants. *International Journal of Systematic and Evolutionary Microbiology*, *63*(Pt_7), 2520-2525. https://doi.org/10.1099/ijs.0.046011-0

Navarrete, I. A., Gabiana, C. C., Dumo, J. R. E., Salmo, S. G., Guzman, M. A. L. G., Valera, N. S., & Espiritu, E. Q. (2017). Heavy metal concentrations in soils and vegetation in urban areas of Quezon City, Philippines. *Environmental Monitoring and Assessment*, *189*(4), 1-15. https://doi.org/10.1007/s10661-017-5849-y

Pedersen, J. S., Carstens, A. B., Djurhuus, A. M., Kot, W., Neve, H., & Hansen, L. H. (2020). *Pectobacterium* phage Jarilo displays broad host range and represents a novel genus of bacteriophages within the family *Autographiviridae*. *PHAGE*, *1*(4), 237-244. https://doi.org/10.1089/phage.2020.0037

Pérombelon, M. C. M. (2002). Potato diseases caused by soft rot erwinias: an overview of pathogenesis. *Plant Pathology*, *51*(1), 1-12. https://doi.org/10.1046/j.0032-0862.2001.Shorttitle.doc.x

Petriccione, M., Zampella, L., Mastrobuoni, F., & Scortichini, M. (2017). Occurrence of copper-resistant *Pseudomonas syringae* pv. *syringae* strains isolated from rain and kiwifruit orchards also infected by *P. s.* pv. *actinidiae*. *European Journal of Plant Pathology*, *149*(4), 953-968. https://doi.org/10.1007/s10658-017-1246-1

Pollumaa, L., Alamae, T., & Mae, A. (2012). Quorum sensing and expression of virulence in pectobacteria. *Sensors (Basel)*, *12*(3), 3327-3349. https://doi.org/10.3390/s120303327

Portier, P., Pédron, J., Taghouti, G., Fischer-Le Saux, M., Caullireau, E., Bertrand, C., Laurent, A., Chawki, K., Oulgazi, S., Moumni, M., Andrivon, D., Dutrieux, C., Faure, D., Hélias, V., & Barny, M.-A. (2019). Elevation of *Pectobacterium carotovorum* subsp. *odoriferum* to species level as *Pectobacterium odoriferum* sp. nov., proposal of *Pectobacterium brasiliense* sp. nov. and *Pectobacterium actinidiae* sp. nov., emended description of *Pectobacterium carotovorum* and description of *Pectobacterium versatile* sp. nov., isolated from streams and symptoms on diverse plants. *International Journal of Systematic and Evolutionary Microbiology*, *69*(10), 3207-3216. https://doi.org/10.1099/ijsem.0.003611

Rahman, M. M., Ali, M. E., Khan, A. A., Akanda, A. M., Uddin, M. K., Hashim, U., & Abd Hamid, S. B. (2012). Isolation, characterization, and identification of biological control agent for potato soft rot in Bangladesh. *The Scientific World Journal*, *2012*, 1-6. https://doi.org/10.1100/2012/723293

Roach, R., Mann, R., Gambley, C. G., Shivas, R. G., Chapman, T., & Rodoni, B. (2020). Pathogenicity and copper tolerance in Australian *Xanthomonas* species associated with bacterial leaf spot. *Crop Protection*, *127*(104923), 1-11. https://doi.org/10.1016/j.cropro.2019.104923

Ross, A., Ward, S., & Hyman, P. (2016). More is better: selecting for broad host range bacteriophages. *Frontiers in Microbiology*, *7*(SEP), 1-6. https://doi.org/10.3389/fmicb.2016.01352

Shneider, M. M., Lukianova, A. A., Evseev, P. V., Shpirt, A. M., Kabilov, M. R., Tokmakova, A. D., Miroshnikov, K. K., Obraztsova, E. A., Baturina, O. A., Shashkov, A. S., Ignatov, A. N., Knirel, Y. A., & Miroshnikov, K. A. (2020). *Autographivirinae* bacteriophage Arno 160 infects *Pectobacterium carotovorum* via depolymerization of the bacterial O-polysaccharide. *International Journal of Molecular Sciences*, *21*(9), 3170-3170. https://doi.org/10.3390/ijms21093170

Smolarska, A., Rabalski, L., Narajczyk, M., & Czajkowski, R. (2018). Isolation and phenotypic and morphological characterization of the first *Podoviridae* lytic bacteriophages ☐A38 and ☐A41 infecting *Pectobacterium parmentieri* (former *Pectobacterium wasabiae*). *European Journal of Plant Pathology*, *150*(2), 413-425. https://doi.org/10.1007/s10658-017-1289-3

Sulakvelidze, A., Alavidze, Z., & Morris, J. G. (2001). Bacteriophage therapy. *Antimicrobial Agents and Chemotherapy*, *45*(3), 649-659. https://doi.org/10.1128/AAC.45.3.649-659.2001

Sundin, G. W., & Wang, N. (2018). Antibiotic resistance in plant-pathogenic bacteria. *Annual Review of Phytopathology*, *56*(1), 161-180. https://doi.org/10.1146/annurev-phyto-080417-045946

Thomas, E. (2001). Multiplicity of infection (*Encyclopedia of Genetics* (pp. 1258-1258). Elsevier. https://doi.org/10.1006/rwgn.2001.0858

Voronina, M. V., Bugaeva, E. N., Vasiliev, D. M., Kabanova, A. P., Barannik, A. P., Shneider, M. M., Kulikov, E. E., Korzhenkov, A. A., Toschakov, S. V., Ignatov, A. N., & Miroshnikov, K. A. (2019). Characterization of *Pectobacterium carotovorum* subsp. *carotovorum* bacteriophage PP16 prospective for biocontrol of potato soft rot.

Microbiology (Russian Federation), *88*(4), 451-460.
https://doi.org/10.1134/S0026261719040118
Wallace, G. B. (1948). Bacterial soft rot of vegetables. *The East African Agricultural*
Journal, *14*(1), 34-35. https://doi.org/10.1080/03670074.1948.11664644
Zaczek-Moczydłowska, M. A., Young, G. K., Trudgett, J., Plahe, C., Fleming, C. C.,
Campbell, K., & O'Hanlon, R. (2020). Phage cocktail containing *Podoviridae* and
Myoviridae bacteriophages inhibits the growth of *Pectobacterium* spp. under in vitro
and in vivo conditions. *PLOS ONE*, *15*(4), e0230842.
https://doi.org/10.1371/journal.pone.0230842
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759 **Tables and Figures**

760 **Table 1.** Characteristics and biological properties of the three lytic bacteriophages isolated using *Pectobacterium carotovorum* subsp.

761 *carotovorum.*

Phage	Morphology	Head diameter (nm)	Tail Length (nm)	Plaque Morphology		Optimal MOI	Latent Period (min)	Burst Size (PFU/cell)
				Transparency	Diameter (mm)			
vB_PcaP-A3	Podovirus	81.07 ± 7.04	19.58 ± 2.92	clear	2.0 - 3.0	0.01	10	142
vB_PcaM-D1	Myovirus	77.50 ± 4.68	98.15 ± 4.09	clear	1.0 - 1.9	0.1	10	151
vB_PcaM-J3	Myovirus	76.17 ± 2.81	137.08 ± 12.78	clear	< 1.0	10	30	42

762   **Figure 1. Transmission electron micrographs (TEM) of vB_PcaP-A3, vB_PcaM-D1, and**

763   **vB_PcaM-J3 virions.** High titer phage lysates were negatively stained with 2% aqueous

764   uranyl acetate and were visualized by transmission electron microscopy (TEM; JEOL JEM-

765   1220 Electron Microscope). The images were scanned with 800 ms exposure, 1.4 gain, and

766   1.0 bin with no sharpening, normal contrast, and 1.00 gamma. Bar represents 50nm.

767 **Figure 2**. **Growth characteristics of phages vB_PcaP-A3, vB_PcaM-D1, and vB_PcaM-**

768 **J3 and their stability on varying temperature and pH levels.** **(A)** One-step growth curve

769 of each phage grown together with log phase *Pectobacterium carotovorum* subsp.

770 *carotovorum* in TSB for 60 min at 30 °C. **(B)** Adsorption of each phage to log phase

771 *Pectobacterium carotovorum* subsp. *carotovorum* in TSB for 60 min at 30 °C. **(C)** Stability

772 of the phages at various temperature and pH levels.

Figure 3. Bacterial killing curves of phages vB_PcaP-A3, vB_PcaM-D1, and vB_PcaM-

J3 and their protective effect against *Pectobacterium carotovorum* subsp. *carotovorum* on

potato tuber slices based on optimal MOI. (A) Killing curves of phages vB_PcaP-A3,

vB_PcaM-D1, vB_PcaM-J3 against *Pectobacterium carotovorum* subsp. *carotovorum*

measured as change in absorbance at $OD_{600 \text{ nm}}$. **(B)** Protective effect of monophage treatment

on potato tuber slices against *Pectobacterium carotovorum* subsp. *carotovorum*. **(C)** Mean

diameter of rotting tissue. **(D)** Percent tissue maceration reduction in potato slices co-

inoculated with *Pectobacterium carotovorum* subsp. *carotovorum* and vB_PcaP-A3,

vB_PcaM-D1 or vB_PcaM-J3 and **(E)** the preventive-curative effect of selected

bacteriophages in potato tubers against *Pectobacterium carotovorum* subs. *carotovorum* after

72-h incubation at 28°C. Each dot in **(C-D)** is an individual potato slice (ie. replicate).

Statistical significance was determined with one-way ANOVA with Tukey's post-test and

one-sample t-test in **(C)** and **(D)**, respectively, ****$p < 0.0001$.

Figure 4. Suppression of *Pectobecterium carotovorum* subsp. *carotovorum* by phage

cocktails *in vitro* and semi-*in planta*. (A) Lytic activities of cocktails of two (vB_PcaP-

A3+vB_PcaM-D1, vB_PcaP-A3+vB_PcaM-J3, and vB_PcaM-D1+vB_PcaM-J3) and three

(vB_PcaP-A3+vB_PcaM-D1+vB_PcaM-J3) phages at low and high MOI against

Pectobacterium carotovorum subsp. *carotovorum* measured as change in absorbance at

$OD_{600 \ nm}$. **(B)** Protective effect of the phage cocktails on potato tuber slices against

Pectobacterium carotovorum subsp. *carotovorum*. **(C)** Mean diameter of rotting tissue and

(D) percent tissue maceration reduction in potato slices co-inoculated with *Pectobacterium*

carotovorum subsp. *carotovorum* and phage cocktails.

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