

2021 Third Year Phage Hunters

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Welcome Phage Hunters!

Introduction

Lab times

Lab sessions will run from 0930-1230 each day in GP108. The kick-off meeting (Monday 17th May) where students will be handed a sampling pack will take place in a lecture theatre to be determined.

Chapter 1

Phage Enrichment

In total, we will have 15 students isolating phages for the CPL. Each student will be assigned a pathogen and will use that to isolate phages from samples crowdsourced by their peers. The first week is spent enriching for phages from their samples. In total there will be 47 samples (plus one negative ctrl).

1.1 Monday 17th May

- Kick off meeting + students given sample packs, with samples to be taken next morning.

BT will give a presentation on the work and the steps involved in phage isolation.

1.1.1 Materials required

- 36 sample jars
- 12 lab pens

1.2 Tuesday 18th May

- Students return to lab with samples
- Samples are transferred to 50 mL falcon tube + centrifuged for 30 mins @ 10000 x g
- Samples are filtered through 0.2 μm syringe filters into fresh 50 mL Falcon tube, then aliquoted into 15 1.5 mL lo-bind microcentrifuge tubes
- Samples are recorded on class shared spreadsheet
- Samples stored O/N at 4 °C.

1.2.1 Materials required

- 6 x 50 mL Falcon tube x 15 students (90 total)
- 3 x 25 mL syringe with luer lock (45 total)
- 3 x 0.2 μm syringe filter (45 total)
- 3 x 15 1.5 mL microcentrifuge tubes (675 total)
- microcentrifuge rack
- Centrifuge capable of 10,000 x g for 50 mL Falcon tubes

1.3 Wednesday 19th May

- Students assigned a pathogen
- Students provided a set of samples
- Students aliquot 900 μL of each sample into each well of a 96-well deep well plate
- Students aliquot 500 μL of 3x LB + 30 mM MgCl_2 + 30 mM CaCl_2 into each well
- Students aliquot 100 μL of O/N pathogen culture into each well.
- Cover and incubate overnight at 30 °C on an orbital shaker.
- Students set up fresh O/N culture of pathogen

1.3.1 Materials required

- 10 mL of O/N pathogen culture
- 50 mL of 3x LB + 30 mM MgCl_2 + 30 mM CaCl_2 (1L total)
- Deep well plate (15 total)
- PCR film (15 total)
- 2 sterilins with 10 mL LB + 10 mM MgCl_2 + 10 mM CaCl_2 (30 total)

1.4 Thursday 20th May

- Students transfer 200 μL from each well into a 0.2 μm filter plate atop a regular, sterile 96 well plate.
- Plate is spun at 900 x g for 4 mins to transfer filtrate to bottom plate
- Into a fresh 200 μL 96-well plate, students add 190 μL of LB + 10 mM MgCl_2 + 10 mM CaCl_2 per well
- 5 μL of phage lysate from bottom plate is added to each well
- 10 μL of O/N host culture added to each well and plate sealed
- Placed in an incubator O/N at 37 °C.
- Students set up fresh O/N culture of pathogen

1.4.1 Materials required

- 0.45 μm filter plate (Merck MSHAS4510) (15 total)
- 2 x sterile 200 μL 96-well plate (30 total)
- PCR film (15 total)
- 2 sterilins with 10 mL LB + 10 mM MgCl_2 + 10 mM CaCl_2 (30 total)

1.5 Friday 21st May

- Students transfer 200 μL from each well into a 0.2 μm filter plate atop a regular, sterile 96 well plate.
- Plate is spun at 900 x g for 4 mins to transfer filtrate to bottom plate
- Bottom plate is covered with PCR film and stored until Monday at 4 $^{\circ}\text{C}$.
- Students streak pathogen onto agar plate for growth over weekend.

1.5.1 Materials required

- 0.45 μm filter plate (15 total)
- 1 x sterile 200 μL 96-well plate (15 total)
- PCR film (15 total)
- 2 x LB agar plates for bacterial growth
- Inoculating loops

Chapter 2

Spot Assays and Purification

In the second week, the students will be performing spot assays and phage purification.

2.1 Monday 24th May

Students will set up four top agar spot plates, with 12 samples per plate (w. neg. ctrl).

- Students will mark plates into quadrants
- Students will add 1 mL of host culture at OD_{600} of 0.6 to 3 mL of molten top agar (0.6% agar in LB + 10 mM $MgCl_2$ + 10 mM $CaCl_2$), vortex briefly and then pour onto bottom plate
- Once the plates are set, students will spot 5 μ L of phage lysate from each sample at 3 per quadrant
- Plates will be incubated O/N at 37 °C for plaque formation

2.1.1 Materials required

- 10 mL pathogen culture
- 5 glass test tubes with lids containing 3 mL molten agar (75 total)
- 5 petri dishes with bottom agar (75 total)
- water bath (or heat blocks if the tubes will fit in).

2.2 Tuesday 25th May

Students will pick plaques and perform first round of O/N purification steps. We are going to assume a maximum of 3 phages per student.

- Students will take a photo of their plates to record plaque morphology
- For each plaque, students will core the plaque using a pipette tip and place it in 200 μ L of SM buffer in a microcentrifuge tube, before vortexing briefly.
- Students will then set up a dilution series from 10^0 to 10^{-11} in SM Buffer in a 96 well plate (one row per phage)
- Using the same method as the spot assay, students will spot the 12 dilutions onto a single plate (so one plate per phage).
- Plates will be left O/N for plaques to develop

2.2.1 Materials required

- 10 mL pathogen culture
- 3 glass test tubes with lids containing 3 mL molten agar (45 total)
- 3 petri dishes with bottom agar (45 total)
- water bath (or heat blocks if the tubes will fit in).
- 1 sterile 96 well plate (15 total)
- 3 lo-bind microcentrifuge tubes (45 total)

2.3 Wednesday 26th May

From each plate, students will pick plaque from the biggest dilution (fewest phages) and perform second round of O/N purification. We are going to assume a maximum of 3 phages per student.

- Students will take a photo of their plates to record plaque morphology
- For each plaque, students will core the plaque using a pipette tip and place it in 200 μ L of SM buffer in a microcentrifuge tube, before vortexing briefly.
- Students will then set up a dilution series from 10^0 to 10^{-11} in SM Buffer in a 96 well plate (one row per phage)
- Using the same method as the spot assay, students will spot the 12 dilutions onto a single plate (so one plate per phage).
- Plates will be left O/N for plaques to develop

2.3.1 Materials required

- 10 mL pathogen culture
- 3 glass test tubes with lids containing 3 mL molten agar (45 total)
- 3 petri dishes with bottom agar (45 total)
- water bath (or heat blocks if the tubes will fit in).
- 1 sterile 96 well plate (15 total)
- 3 lo-bind microcentrifuge tubes (45 total)

2.4 Thursday 27th May

From each plate, students will pick plaque from the biggest dilution (fewest phages) and perform third round of O/N purification. We are going to assume a maximum of 3 phages per student.

- Students will take a photo of their plates to record plaque morphology
- For each plaque, students will core the plaque using a pipette tip and place it in 200 μ L of SM buffer in a microcentrifuge tube, before vortexing briefly.
- Students will then set up a dilution series from 10^0 to 10^{-11} in SM Buffer in a 96 well plate (one row per phage)
- Using the same method as the spot assay, students will spot the 12 dilutions onto a single plate (so one plate per phage).
- Plates will be left O/N for plaques to develop
- Students will set up O/N cultures of host

2.4.1 Materials required

- 10 mL pathogen culture
- 3 glass test tubes with lids containing 3 mL molten agar (45 total)
- 3 petri dishes with bottom agar (45 total)
- water bath (or heat blocks if the tubes will fit in).
- 1 sterile 96 well plate (15 total)
- 3 lo-bind microcentrifuge tubes (45 total)
- 2 sterilins containing 10 mL LB + 10 mM $MgCl_2$ + 10 mM $CaCl_2$ (30 total)

2.5 Friday 28th May

From each plate, students will pick plaque from the biggest dilution (fewest phages) and bulk it up on hosts overnight.

- Students will take a photo of their plates to record plaque morphology. These will also be used to determine approximate PFU counts.
- Students will pick plaque from the biggest dilution (fewest phages) and add it to 50 mL of LB + 10 mM $MgCl_2$ + 10 mM $CaCl_2$, amended with 1 mL of O/N host culture.
- Cultures will be grown O/N at 30 °C

The next day (Saturday), BT and team will go in and centrifuge the samples down and prepare for each phage 1 x 50 mL tubes of lysate, filtered through a 0.2 μ m syringe filter. We will prepare 3 x 2 mL phage stocks in acid washed, autoclaved amber glass vials.

200 μ L of each phage filtrate will be provided to the imaging centre.

2.5.1 Materials required

- 3 x 50 mL LB + 10 mM $MgCl_2$ + 10 mM $CaCl_2$ (2L total)
- 3 x 50 mL falcon tubes (45 total)
- 3 x 0.2 μ m syringe filter (45 total)
- 3 x 25 mL syringe with luer lock (45 total).
- 3 x 2 mL amber glass vials

Chapter 3

Phage Purification and DNA extraction

In the third week, the students will perform DNA extractions on phages

3.1 Tuesday 1st June

Students will test their lysate for estimating PFU and begin phage DNA precipitation

- Students will prepare a dilution series of their phages as before and spot them in triplicate across 3 plates
- Students will transfer 30 mL of lysate to a fresh Falcon tube
- Students will add 15 μ L of nuclease solution and incubate at 37 °C for 30 mins
- Students will then add 15 mL of precipitant solution to each tube and mix gently by inversion
- Samples will be incubated at 4 °C O/N

3.1.1 Materials required

- 10 mL pathogen culture
- 3 glass test tubes with lids containing 3 mL molten agar (45 total)
- 3 petri dishes with bottom agar (45 total)
- water bath (or heat blocks if the tubes will fit in).
- 1 sterile 96 well plate (15 total)
- 3 x Falcon tube (45 total)

- 50 μL of nuclease solution (750 μL total)
- 50 mL of precipitant solution (750 mL total)

3.2 Wednesday 2nd June

Students will extract DNA using the Promega Wizard kit using this protocol

Depending on how many phages we isolate, we are anticipating sending two per student for sequencing, with the remaining extractions kept back for future analyses. ### Materials required * 3 x 500 μL resuspension buffer (5 mM MgSO_4) (25 mL total) * 2 x Promega Wizard Kits.

3.3 Thursday 3rd June

Students will perform killing efficiency assays of their phages.

Group photo!

3.4 Friday 4th June

Christian and Lauren will provide an overview on using the tools via Zoom.

Two videos from the CPT also exist for this purpose:

- Structural Pipeline Video
- Functional Pipeline Video

Chapter 4

Genome Annotation

The students will use the Centre for Phage Technology Galaxy Phage tool to annotate their genomes

4.1 Monday 7th June

Sequencing data comes back from the sequencing centre. BT performs assembly and read mapping and gives data to the students

4.2 Tuesday 8th - Friday 11th June

Students will embark on analysing their genomes. This will include:

1. Classification using VIPTree
2. Identification of novel species and genera through VIRIDIC

Recipes for reagents

4.3 Precipitant solution

This is a ready-mixed 30% w/v PEG-8000, 3 M NaCl solution for adding to phage lysate in a 1:2 ratio precipitant:lysate (10%PEG-8000, 1 M NaCl final conc.)

1. In a sterilized 500 mL bottle add 330 mL of autoclaved MilliQ and 105 g of NaCl and dissolve.
2. Add 180g of PEG8000, cap bottle and shake.
3. Incubate bottle in a 60 °C waterbath for 3 hours, shaking occasionally
4. Remove and let cool to RT, shaking occasionally
5. Add autoclaved MilliQ to 600 mL and store at RT.

4.4 Beef resuspension solution

This is based on the recipe from (Williamson et al., 2003). For each sample of 25g:

1. To a 500 mL Duran add the following:
 - 25g of beef extract
 - 3.35g of $Na_2HPO_4 \cdot 7H_2O$
 - 0.3g of citric acid
2. Add 250 mL of autoclaved MQ
3. Add a stirrer bar
4. Mix until completely dissolved and adjust pH to 7.2

Protocols

4.5 Preparing phages from environmental samples

Note: Due to the need to centrifuge samples in a fixed rotor, you can only process *eight* samples at any one time

Please read all associated Risk Assessments related to the pathogens of interest before proceeding.

4.5.1 Solid Samples

This method is adapted from (Guzmán et al., 2007). The proteins in the beef extract help phages desorb from solid material. Proteinaceous substances within the beef extract and those that desorb with the phages can interfere with downstream molecular analyses on large samples (e.g. PCR) (Göller et al., 2020), but for phage isolation, this isn't a problem (and beef extract is more cost effective than using PBS + BSA).

1. Aliquot 10-25g of sample into a 250 mL Duran, with a magnetic stirrer
2. Add 1:10 w/v 10% beef resuspension solution (pH 7.2)
3. Homogenise the sample by magnetic stirring for 20 mins at RT with sufficient speed to develop a vortex (500-900 rpm)
4. Transfer to 50 mL falcon tubes or larger vessels and centrifuge at 4080 xg for 10 mins at 4 °C. This stage is just to get rid of the bulk material, so can be done in the benchtop centrifuge in GP211.
5. Pass the supernatant through a Pluriselect 40 µm mesh filter into a fresh 50 mL Falcon tube to the 40 mL mark. Draw an arrow on the lid and mark the side where the pellet is going to be.
6. Centrifuge at 10,000 xg for 30 mins at 4 °C. This requires the ThermoScientific Multifuge X3R on the 4th floor.
7. In a flow hood, filter through a 0.2 µm PES filter into an acid-washed and autoclaved 20 mL amber glass vial (Fisher 11503552). You may have to

re-centrifuge if it's really clogging the filter. Use a luer-lock syringe and filter for this to avoid it popping off and spraying everywhere! You may need to use a second filter if it starts to clog.

8. Label the glass vial with the What3Words label. Also on the side, write the date it was processed and your name.
9. Store at 4 °C in the dark in the CPL sample box

4.5.2 Liquid Samples

1. Aliquot sample into a 50 mL Falcon tube. Draw an arrow on the lid and mark the side where the pellet is going to be.
2. Top up to 45 mL with pH 7.5 buffer if necessary
3. Centrifuge at 10,000 xg for 30 mins at 4 °C. This requires the ThermoScientific Multifuge X3R on the 4th floor.
4. Pour supernatant into 20 mL syringe fitted with 0.2 µm PES filter
5. In a flow hood, filter through a 0.2 µm PES filter into an acid-washed and autoclaved 20 mL amber glass vial (Fisher 11503552). You may have to re-centrifuge if it's really clogging the filter. Use a luer-lock syringe and filter for this to avoid it popping off and spraying everywhere!
6. Label the glass vial with the What3Words label. Also on the side, write the date it was processed and your name.
7. Store at 4 °C in the dark in the CPL sample box

4.6 1. Initial phage enrichment (using 2.2 mL deep well plates or 7 mL sterilins)

This method is based off this work (Olsen et al., 2020) and allows for the screening of one or more host strains against up to 94 samples (plus a blank and a positive control). Briefly, into each well we are adding LB medium into filtered sample that contains phages, then adding in an overnight culture of the pathogen of interest. If there are phages in the sample that infect the host, they will replicate and thus increase in abundance within the well. Subsequent steps then purify these enriched phages.

For a negative control, you want to use a sample that has no phages in it. DI water is recommended for this. You also want to run a positive control (to show that phages can be amplified and plated). For this, you can use a phage lysate that has been previously isolated on the host of interest. Spike 10 μ L of previous phage lysate into 1340 μ L of DI water as your positive control.

If you do not have a phage isolated on the pathogen of interest, it is recommended that you use a positive control consisting of 10 μ L of T4 lysate into 1340 μ L of DI. You will then need a well on your plate that contains an enrichment of *E. coli* BW25113. You'll also need to include this in subsequent plaque assay steps.

1. Make an O/N culture of your host of interest in LB medium at 37 °C on an orbital shaker. The required volume per host is 110 μ L \times the number of samples being tested, plus one negative control and one positive. So for 10 samples plus controls, you will need \sim 2 mL of host culture.
2. Make a 40 mL solution of 0.25 M $CaCl_2$ and 0.25 M $MgCl_2$ as follows:
 - 10 mL 1M $CaCl_2$
 - 10 mL 1M $MgCl_2$
 - 20 mL DI water
 - Vortex and filter sterilize through a 0.22 μ m syringe filter.
3. The next day, unwrap an autoclaved 2.2 ml deep well 96-well plate (make sure you have the right ones! The wells are square. The ones with round wells don't hold enough liquid) in a flow hood.
4. Place 1350 μ L of a 0.45 μ m filtered sample into each well. For every strain of host, you need a negative control well (1350 μ L of DI water). It's worth thinking about the final spot plaque assays when laying out the plate. It's easier to have a whole row or a whole column with a single host strain, as that way, you can spot an entire row/column onto the same plate, rather than having to cherry-pick samples.
5. To each well, add 80 μ L of a mixture of 0.25 M $CaCl_2$ and 0.25 M $MgCl_2$
6. To each well, add in 100 μ L of an O/N culture of host

*4.6. 1. INITIAL PHAGE ENRICHMENT (USING 2.2 ML DEEP WELL PLATES OR 7 ML STERILINS)*²⁷

7. To each well, add in 450 μ L of 4.4x concentration LB (44g of LB in 400 mL)
8. Pipette up and down to mix
9. Cover with a plate film.
10. Incubate overnight at 30 °C on an orbital shaker at 200 rpm.
11. Set up another O/N culture of the hosts for the next day.

Bibliography

- Göller, P. C., Haro-Moreno, J. M., Rodriguez-Valera, F., Loessner, M. J., and Gómez-Sanz, E. (2020). Uncovering a hidden diversity: optimized protocols for the extraction of dsDNA bacteriophages from soil. *Microbiome*, 8(1):17.
- Guzmán, C., Jofre, J., Blanch, A. R., and Lucena, F. (2007). Development of a feasible method to extract somatic coliphages from sludge, soil and treated biowaste. *J. Virol. Methods*, 144(1-2):41–48.
- Olsen, N. S., Hendriksen, N. B., Hansen, L. H., and Kot, W. (2020). A new High-Throughput screening method for phages: Enabling crude isolation and fast identification of diverse phages with therapeutic potential. *PHAGE*.
- Williamson, K. E., Wommack, K. E., and Radosevich, M. (2003). Sampling natural viral communities from soil for culture-independent analyses. *Appl. Environ. Microbiol.*, 69(11):6628–6633.