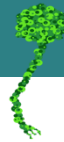


The background features a dark green gradient with several large, semi-transparent circles in shades of green and blue. A stylized phage-like structure, composed of small green circles connected by a thin line, is positioned in the upper center and lower right. A solid teal horizontal band spans the middle of the image, containing the main text.

Phage Genomics Workshop

WELCOME



Karen
Klyczek



Dan
Williams



Steve
Caruso



Veronique
Delesalle



Debbie
Jacobs-Sera



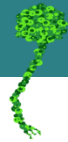
Dan
Russell



Steve
Cresawn



Graham
Hatfull



William
Biederman



Danielle
Heller



Denise
Monti



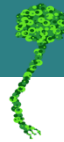
Pushpa
Ramakrishna



Vic
Sivanathan



Bethany
Wise



Expectations for HHMI Events



HHMI welcomes the participation of persons of **all identities** and **backgrounds**.



If you experience or witness **inappropriate behavior**, please speak to your event organizer.



Attendees are expected to treat others with **courtesy** and **respect**.



HHMI will **investigate promptly** and take any appropriate corrective action.



Visit [hhmi.org/policies](https://www.hhmi.org/policies) for details. Refer to your **event materials** for event organizer information.



Event Organizers

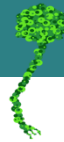


Billy Biederman
biederma@hhmi.org

or Denise Monti
montid@hhmi.org



Vic Sivanathan
sivanathanv@hhmi.org

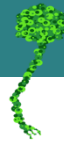


Workshop Logistics

- Keep to workshop schedule (especially for meals)
- Enjoy our campus
- Please wear your name badges



SEA-PHAGES Bioinformatics Workshop Overview



Objectives

- **Characterize and investigate phage genomes** including the details of gene calling, functional assignments, and preparing final submission files **that meet QC requirements**.
 - This year's genome is *Arthrobacter* phage QuinnAvery.
 - **Install and use the software** for annotation and analysis of phages. The software includes DNA Master, Phamerator, PECAAN and other web-based tools.
- **Identify basic phage biology concepts** that underlie bioinformatic investigations.
- **Explore classroom implementation strategies.**



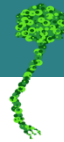
Primary Learning Method

- Learn by doing
 - Gather data
 - Do the work: annotate & analyze the genome
 - Do science like scientists do it. [Translation: new technology changes at a moment's notice]



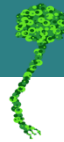
Workshop Logistics

- Workshop page at seaphages.org
- We will be working in groups (assignments are on Workshop page)
- SEA-PHAGES Bioinformatics Guide and the new SEA-PHAGES Genomics Guide
- Lab notebook. Take notes while annotating!
- seaphages.org forums: Check often.



We Will...

- Use *Arthrobacter* phages QuinnAvery genome to
 - Learn to manipulate the tools we use
 - DNA Master
 - GeneMark, Glimmer
 - BLAST
 - Phamerator
 - HHpred
 - Starterator
 - PECAAN
 - Practice calling genes and functions
- Increase our phage biology knowledge
- Produce a **quality** genome annotation
 - Sequence quality control: ✓
 - Annotation quality control: Checklist and files



SEA-PHAGES is an authentic CRE as part of an iREC

Phage Discovery

September 1

Phage Isolation, Purification, Amplification

Phage Characterization, Archiving, DNA Extraction

Sequencing, Assembly, QC

Between semesters

Phage Genomics

Phage Genome Annotation

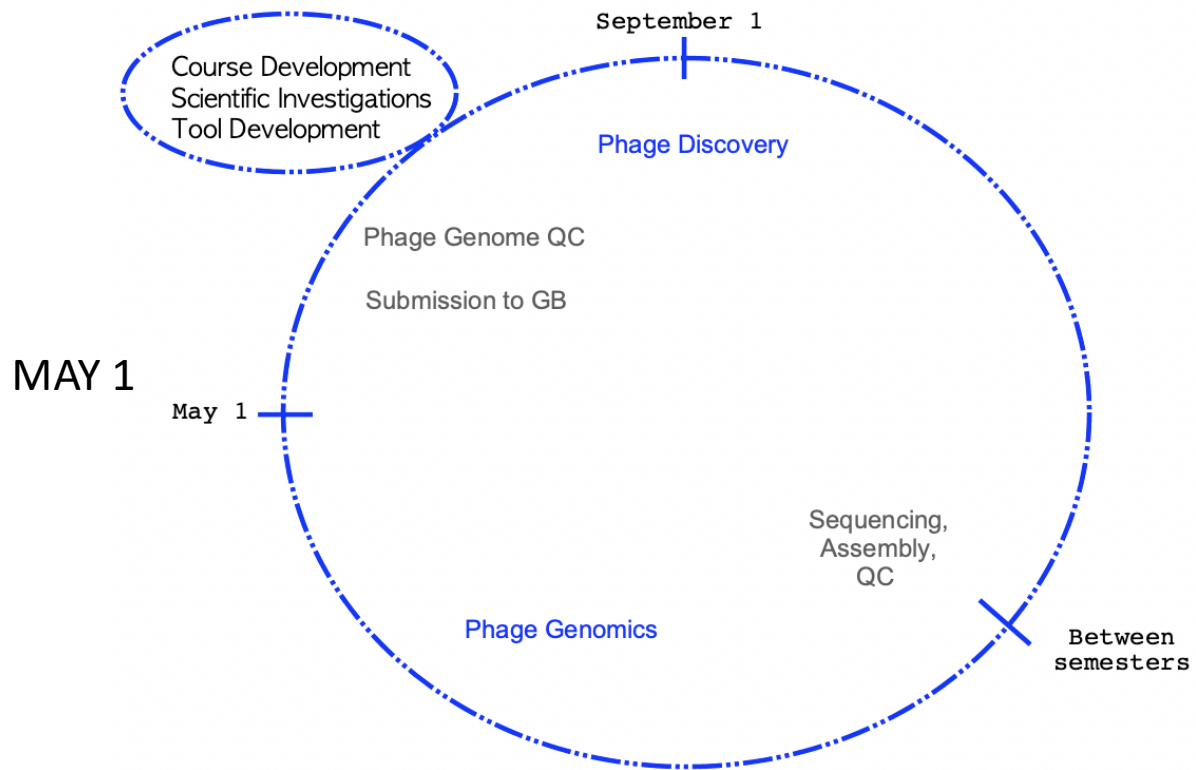
Phage Genome QC

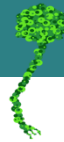
Submission to GB

May 1



September 1





Phage Genomics

Phage Genome Annotation

due May 1

What does Phage Genome Annotation include?

Preliminary Genome Investigation

Zoom out: Whole genome, Whole genome comparisons

Zoom in: Synteny, Gene Functions

Manual Genome/Gene Investigation

3 Questions:

Is it a gene?

What is its start?

What is its function?

Tools:

Auto-Annotation

Phamerator

GeneMark Output

Starterator

Blast

HHPred:

Database Comparisons

PDB

UniProt

CDD

InterPro

Mechanics:

Add a gene

Delete a gene

Change a start

File Preparation

Review/Revise

Document findings

Big picture: zoom out

Whole genome

Cluster, Subcluster evaluation

Comparison decisions

Details: zoom in

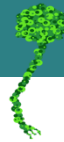
DeNovo decisions

Single genes

Start sites

Gene functions

Special features



Two aspects that go hand in hand: Biology of phages & Mechanics of phage annotation

- To get started on the Biology of phages, review Central Dogma principles, according to phages. (before dinner)
 - Read [Bioinformatics Section](#) in the Phage Discovery Guide
 - Review Central Dogma check-list
 - Questions
- To get started on the Mechanics. (after dinner)
 - Auto-annotate
 - Blast
 - Gather GeneMark graphic outputs
 - Bookmark websites