

A collaborative research effort between the Parfrey Lab and IMERSS

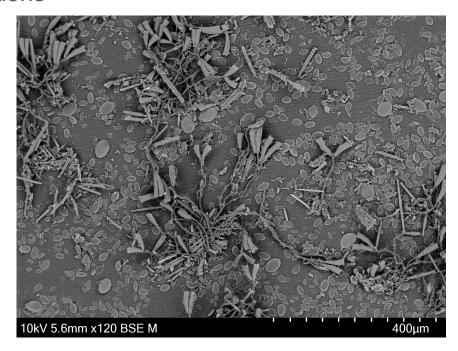
Research objectives meeting 21 October 2020

Background and motivation for project

- Microbial communities (bacteria and protists) on eelgrass leaves are distinct from the water column (Crump et al. 2018; Segovia et al. accepted)
- Bacterial communities vary spatially; different meadows host different communities (Bengtsson et al. 2017; Fahimipour et al. 2017)
- Bacterial community composition varies based on leaf age (Segovia et al. in prep)
- Eelgrass hosts a core microbiome; some taxa are always present on eelgrass leaves even after a disturbance (Adamczyk et al. in prep)
- Diatom communities vary throughout eelgrass leaf; different sections host different communities (Webber and van Asselt preliminary results)

Project goals

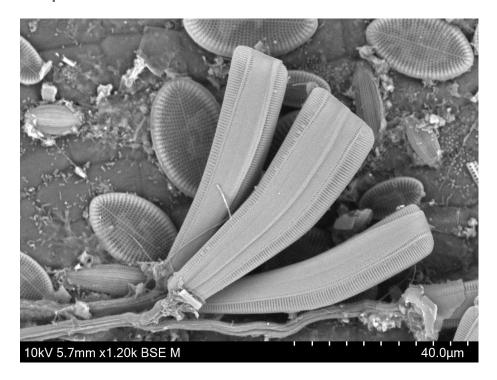
- Determine if eelgrass leaves host different taxa on different sections of the leaf
- Understand bacteria X diatom interactions
 - Members and community structure



Overarching question

Are there consistent interactions between specific bacteria and diatom taxa on

certain sections of eelgrass leaves?



Sub-questions

- 1a) What is the spatial structure of the eelgrass leaf epiphyte community?
 - Across the same leaf section (tip vs tip)
 - Across different leaf section (tip vs middle vs base)
- 1b) Where are these groups found on eelgrass leaves?
 - Diatoms
 - Bacteria
 - Fungi?
- 2) Is the eelgrass diatom community distinct from the water column diatom community?
- 3) Is the epiphyte community different on the edge vs the middle of the meadow?

Proposed general methods and workflow

November sampling -> method validation and data preliminary analysis

Project timeline

Nov. 2020 Preliminary field work

Spring 2020

Succession field work

Sequencing budget

| Sample type | Number of samples needed | Sequencing cost per sample | Total cost for sample type |
|-----------------------|--|----------------------------|----------------------------|
| Eelgrass bacteria | 30 (3 sections from 10 different leaves) | ~\$35 | ~\$1050 |
| Eelgrass diatoms | 30 (3 sections from 10 different leaves) | ~\$35 | ~\$1050 |
| Water column bacteria | 6 (3 within meadow, 3 outside meadow) | ~\$35 | ~\$210 |
| Water column diatoms | 6 (3 within meadow, 3 outside meadow) | ~\$35 | ~\$210 |
| | | TOTAL | ~\$2520 |

SEM and total budget

| Sample type | Number of samples needed | SEM cost per sample | Total cost per sample type |
|----------------------------|--|---------------------|----------------------------|
| Eelgrass | 30 (3 sections from 10 different leaves) | \$28 | \$840 |
| Eelgrass (prelim sampling) | 3? | \$28 | \$84 |

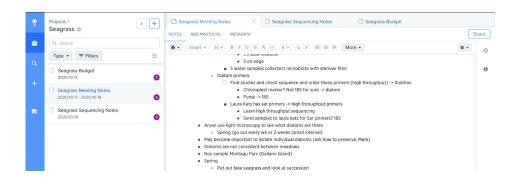
| Sequencing | ~\$2520 |
|------------|---------|
| SEM | ~\$840 |
| TOTAL | ~\$3360 |

Roles

- Siobhan Schenk (UBC) project lead
- Emily Adamczyk (UBC/IMERSS) project manager/advisor
- Elaine Humphrey (UVic/IMERSS) SEM
- Arjan van Asselt (IMERSS) microscopy
- Mark Webber (IMERSS) microscopy
- Laura Parfrey (UBC) project advisor
- Laura Katz? (Smith College) collaborator

Standards for data and collaboration

Benchling e-notebook



- Communication
 - 1 work week to answer emails*
 - Ask for help when needed
- Data sharing and storage
 - Check in before giving anyone access to raw data/records
 - Maintain records according to UBC guidelines *(uVic for Elaine)