

Molecular assessment of wild *Achatinella mustelina* diet

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Food Similarity Between Proposed Donor and Enclosure Snail Sites

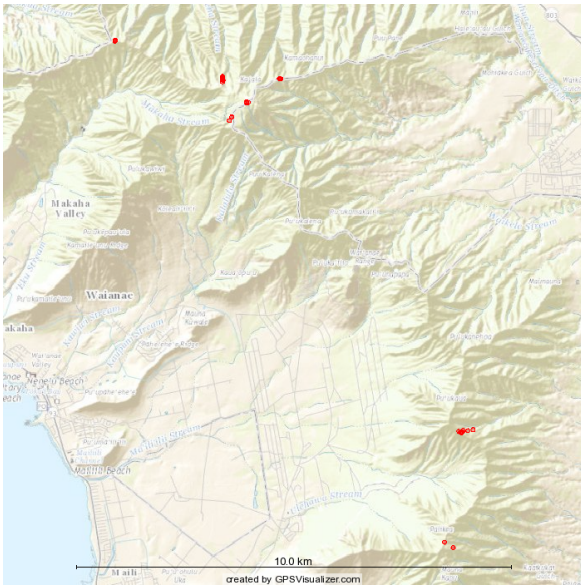


Figure 1 - Map of sampling locations (Waianae range)

If populations of *Achatinella mustelina* in difficult-to-access areas are to be successfully relocated to enclosures at sites more amenable to conservation efforts, it must be assured that conditions at the proposed sites are similar to those where the snails currently reside. One factor that may be important is the availability of preferred snail food sources. We are determining whether epiphytic microbial communities are similar between donor and proposed enclosure sites by sequencing DNA amplicons of material swabbed from the surface of leaves at each location.

At each current and proposed snail site, leaves from at least 10 plants were recorded, collected and returned to the lab. In the lab, leaf surfaces were swabbed and these swabs were subjected to DNA sequencing to determine species composition. If leaf-surface microbial communities are similar between current and proposed sites, it is an indication that food source and availability will not be limiting factors in snail health at proposed sites following translocation. If microbial communities are dissimilar, further work will be done to determine whether these differences are functionally meaningful and/or whether it is possible to inoculate plant surfaces at the proposed sites with microbial food sources from the current sites to ease any potential snail relocation shock.

Donor Site	Proposed Site
Skeet Pass	Ka'ala Bog
Culvert 69	Three Points
Ekahanui	Palikea Area

Fungal ITS genetic marker regions were amplified and sequenced at the University of Hawaii HIMB Genetics Core Facility. The resulting reads were then

binned into probable operational taxonomic units which were used to construct dissimilarity measures between sites.

Results

There was a significant difference (Anosim; $P=0.001$) in epiphyte fungal community structure between the four sites near Kaala summit (Kaala Bog, Culvert 69, Skeet Pass, and Three-Points) and the three sites near Palikea (DS Palikea, Kaaikukai, and Ekahanui). There was no detectable difference between any of the current/proposed site pairings, however (See Fig. 3). Thus, it does not seem likely that any snails moved to proposed enclosure locations will encounter significantly different food sources from their currently paired extant sites and food sources will not be limiting factors in snail health following translocation.

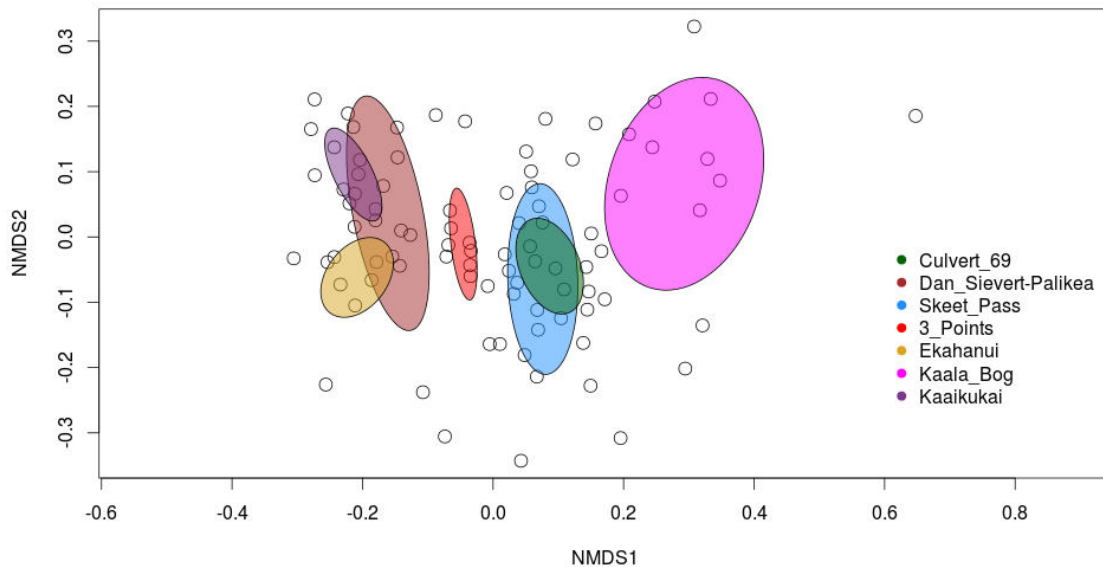


Figure 2 - NMDS projection of epiphyte fungal communities. Ellipses represent the standard deviation of point scores around the group centroid.

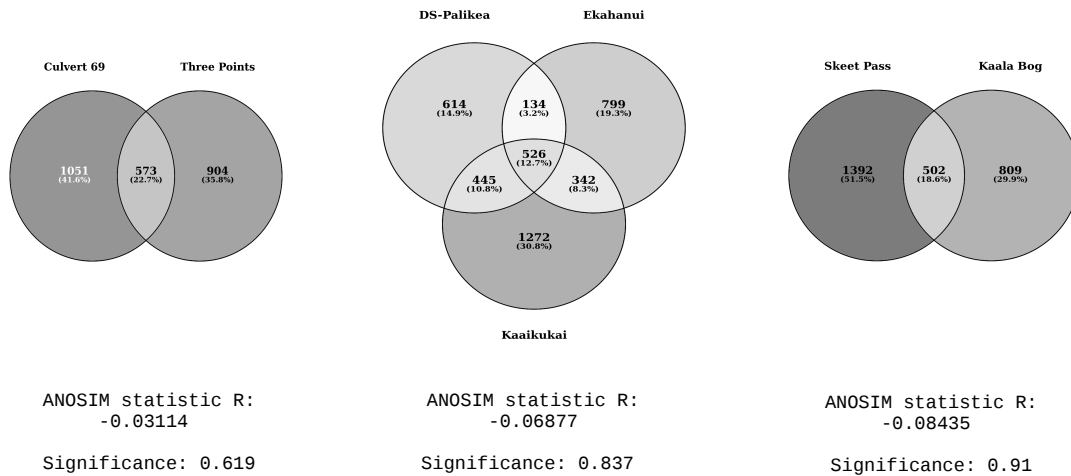


Figure 2 - Shared OTUs between current/proposed site pairings. The proportions of OTUs unique to each site do not constitute statistically significant differences

***Phyllostegia* endophytes and pathogen resistance**

Phyllostegia mollis and *Phyllostegia kaalaensis* are federally listed endangered plant species endemic to Oahu, HI. There are currently no known wild populations of *P. kaalaensis* and the few wild populations of *P. mollis* are failing to demonstrate long-term survival. Greenhouse populations of these plants are maintained by the Army Natural Resources division, but they show marked susceptibility to fungal pathogens, particularly the powdery mildew, *Neoverysiphia galeopsidis*. Greenhouse populations are, therefore, dependent on regular fungicide treatments which are impossible to maintain once individuals have been out-planted to habitats within their native ranges.

Current scientific consensus is that the fungi which coexist within plant tissues form an integral part of plant fitness. These beneficial endophytic and mycorrhizal fungi are not present in plants that have received regular fungicidal treatments, so they are not present in out-planted populations of *P. mollis* or *P. kaalaensis*. One of the major benefits that host plants receive from mutualistic fungi is increased resistance to disease, as mutualistic fungi can outcompete pathogens for habitable living space or even actively repel invasive fungi through excreting chemical compounds. The essentially sterile plants are presumed, therefore, to be highly susceptible to attack by pathogenic fungi in the environment.

We have completed a pilot study on the efficacy of transplanting fungal endophytes from healthy wild populations of *P. mollis* and *P. hirsute* into

greenhouse-raised *P. mollis* and *P. kaalaensis* individuals. Two experimental transplantation methods were tested: 1) Isolating individual fungal strains from wild hosts, culturing them in the lab, and spraying them onto the leaves of greenhouse-raised individuals; 2) Preparing a low-tech slurry from leaves of wild individuals, filtering out large particles, and spraying this onto the new host plants. The first method has the benefit that we know exactly what we are applying to the new host leaves, the second method has the benefit of potentially passing on beneficial fungi that are not amenable to laboratory culture.

Preliminary results were intriguing. The cultured fungal isolates did not appear to confer any advantage over the control group with respect to disease severity, but the group receiving the slurry of wild leaves showed delayed mortality and decreased disease severity for a time (Fig. 4). By the end of one month all plants had generally succumbed to *N. galeopsidis* but the “leaf slurry” treatment warrants further investigation, as it showed some benefit, at least for the first three weeks. By this time the pathogen load on the other two treatments was essentially 100%, with all leaf surfaces covered with sporulating fungus and the slurry-treated plants, in such close proximity, did not last long after. We are nearing the end of a second round of tests, and the results are similar and even more pronounced.

DNA from the inoculae and the initial plant endophyte loads was sequenced and the results are surprising. Roughly 90% of the fungal reads from the leaf slurry treatment, which is showing so much promise, come from *N. galeopsidis*, the same pathogen that appears to be killing the plants (Fig. 5). Leaf samples have been taken at regular intervals during both rounds of testing to track the colonization of plant tissues by fungal inoculae. When these samples are sequenced it will be clearer what fungi were able to establish in the plants, and whether the plants treated with the leaf slurry have been colonized by any strains of *N. galeopsidis*.

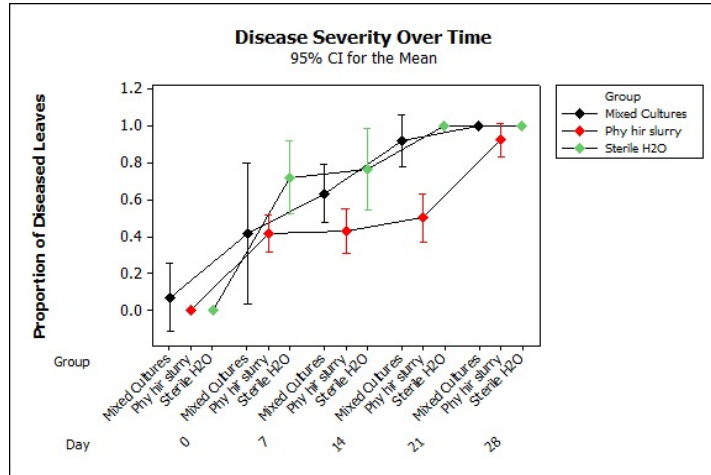


Figure 4 - The proportion of diseased leaves over time. The slurry from healthy wild leaves (shown in red) conferred a longer time until full onset of disease.

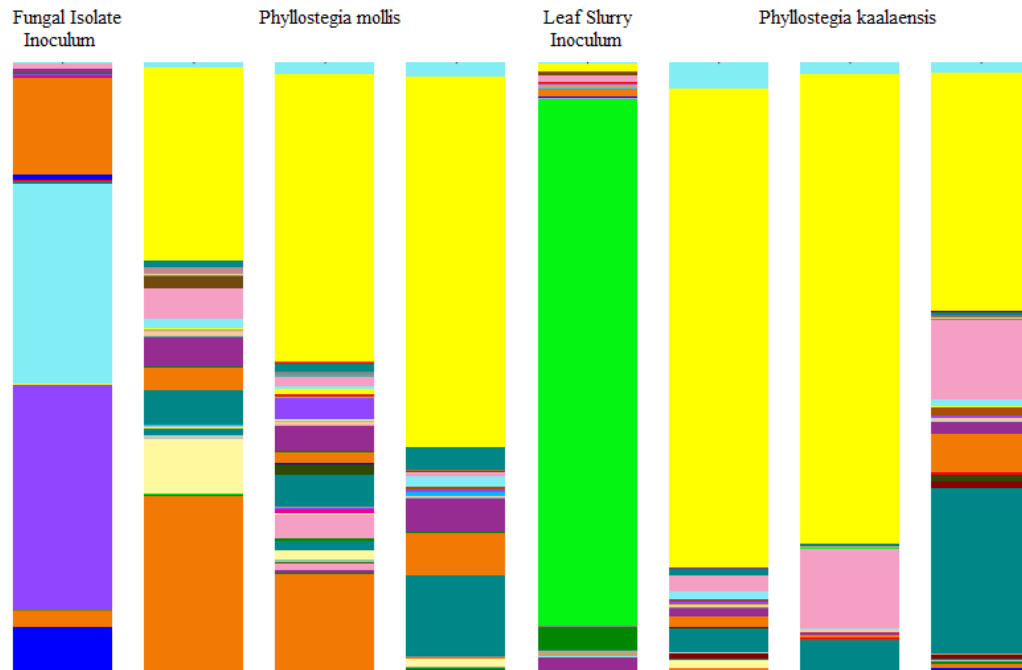


Figure 5 - Stacked bar chart of fungal species identities in initial starting conditions of *Phyllostegia* experiment. The two inoculae, and three replicates of each plant species. The bright green bar in the Leaf Slurry Inoculum represents *N. galeopsidis*.

Other work

Captive (laboratory) snail (*Achatinella mustelina*) populations are dependent on microbes from wild leaves to supplement their diet. These leaves are obtained by regular field forays which are costly and time consuming. The ability to grow diverse microbial communities on laboratory-amenable plants would be a major convenience for maintaining healthy laboratory snail populations.

We are in the early stages of investigating the efficacy of such a system using the model plant *Arabidopsis thaliana*. An initial study is under way to examine the factors that determine the composition of a newly-forming microbial community, such as would be seeded onto the plants in order to grow “snail food.” This plant is very fast growing and there are thousands of curated ecotypes that display a wide range of phenotypic traits, so it potentially offers a highly customizable “delivery system” for supplementing snail captive diets without constant trips into the field.