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Composition change of the phyllosphere due to secondary metabolite produciton

The plant phyllosphere is the culmination of all bacteria, and phages, that are present in the above ground tissues of plants starting at the crown. These communities are comprised of harmful bacterial (such as pathogenic bacteria like *Xanthomonas campestris*), neutral bacteria, and beneficial bacteria. Unlike the root microbiome, there is a lack of nitrogen fixing bacteria such as *Rhyzobia* (which is responsible for the root nodule formation on legumes), and other root microbiome specific bacteria, which cannot successfully grow on the upper portions of plants (1). While the root microbiome has an extensive list of research focused on elucidating the community composition and the interactions between the host-plant and the root microbiome, there is a surprising lack of research looking at the composition and interactions of the phyllosphere with its host plant. Studies looking at the interactions between the host-plant and the root microbiota have given way to complex interactions where the host-plant (*Arabidopsis thailiana*) releases a defence compound (secondary metabolite) called galactinol which is in-turn is utilized as a nutrient by the bacterial pathogen *Agrobacterium fabrum* thereforenegating its anti-microbial effectswhich allows it to resist the host-plants defence response (2). With this in mind, the next question to be asked is if the community of microbiome of the phyllosphere is affected by the release of these defence chemicals. As this is just one of the hundreds of defence chemicals released by the plant for defence there are many unexplored chemicals which may be utilized by organisms in the microbiome for their own use. What this project wishes to investigate is to see the effects that defence compounds (such as Galactinol and Coronatine) have on the composition of the microbiome. It is predicted that chemicals like galactinol may give way to an increase in the presence of certain microbial species within the phyllosphere. Through a mixture of sterile planting of *Arabidopsis thailiana* and controlled application of defense compounds to the leaves of the plant we hope to induce a change in the phyllosphere microbiome. Coupling this with current advances in community composition analysis via Droplet Digital PCR can allow for absolute abundances of bacteria to be gleamed from leaves which will allow for more robust community composition (3).

- The secondary metabolites which are to be investigated are;

Coronatine: A Jasmonic Acid analog which pathogens secrete in order to stop the pathogen response in *A thailiana.*

*Chosen due to it being a chemical released by pathogens in order to circumvent the host- plant from eradicating the pathogen – which may have an effect on the microbiome*

Urate: A common anti-microbial defence compound

*A common defence compound.*

3-(Isothiocyanatomethyl)-1H-indole [ITC]: A glucosinolate compound which is an anti-herbivore agent.

*One of more common mustard herbivore defence compounds. Will be interesting to see the effects it may have on the microbiome.*

Galactinol: A sugar based defence chemical with anti-microbial properties.

*Already known to be utilized by one pathogen as a nutrient and may also be utilized by other microbes as such.*

References

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