

University of Louisville

ThinkIR: The University of Louisville's Institutional Repository

Undergraduate Arts and Research Showcase

Undergraduate Research

2021

The Study of PPAL and its Role in the Development of *Physcomitrella patens*

Susana Perez-Martinez
University of Louisville

Christine Chen
christine.chen@louisville.edu

Mark P. Running
University of Louisville

Follow this and additional works at: <https://ir.library.louisville.edu/uars>



Part of the [Molecular Genetics Commons](#), [Plant Biology Commons](#), and the [Plant Breeding and Genetics Commons](#)

Recommended Citation

Perez-Martinez, Susana; Chen, Christine; and Running, Mark P., "The Study of PPAL and its Role in the Development of *Physcomitrella patens*" (2021). *Undergraduate Arts and Research Showcase*. 60.
<https://ir.library.louisville.edu/uars/60>

This Book is brought to you for free and open access by the Undergraduate Research at ThinkIR: The University of Louisville's Institutional Repository. It has been accepted for inclusion in Undergraduate Arts and Research Showcase by an authorized administrator of ThinkIR: The University of Louisville's Institutional Repository. For more information, please contact thinkir@louisville.edu.

The Study of PPAL and its Role in the Development of *Physcomitrella patens*

Susana Perez Martinez, Christine Chen, Mark P. Running

Department of Biology, University of Louisville, Louisville KY 40292

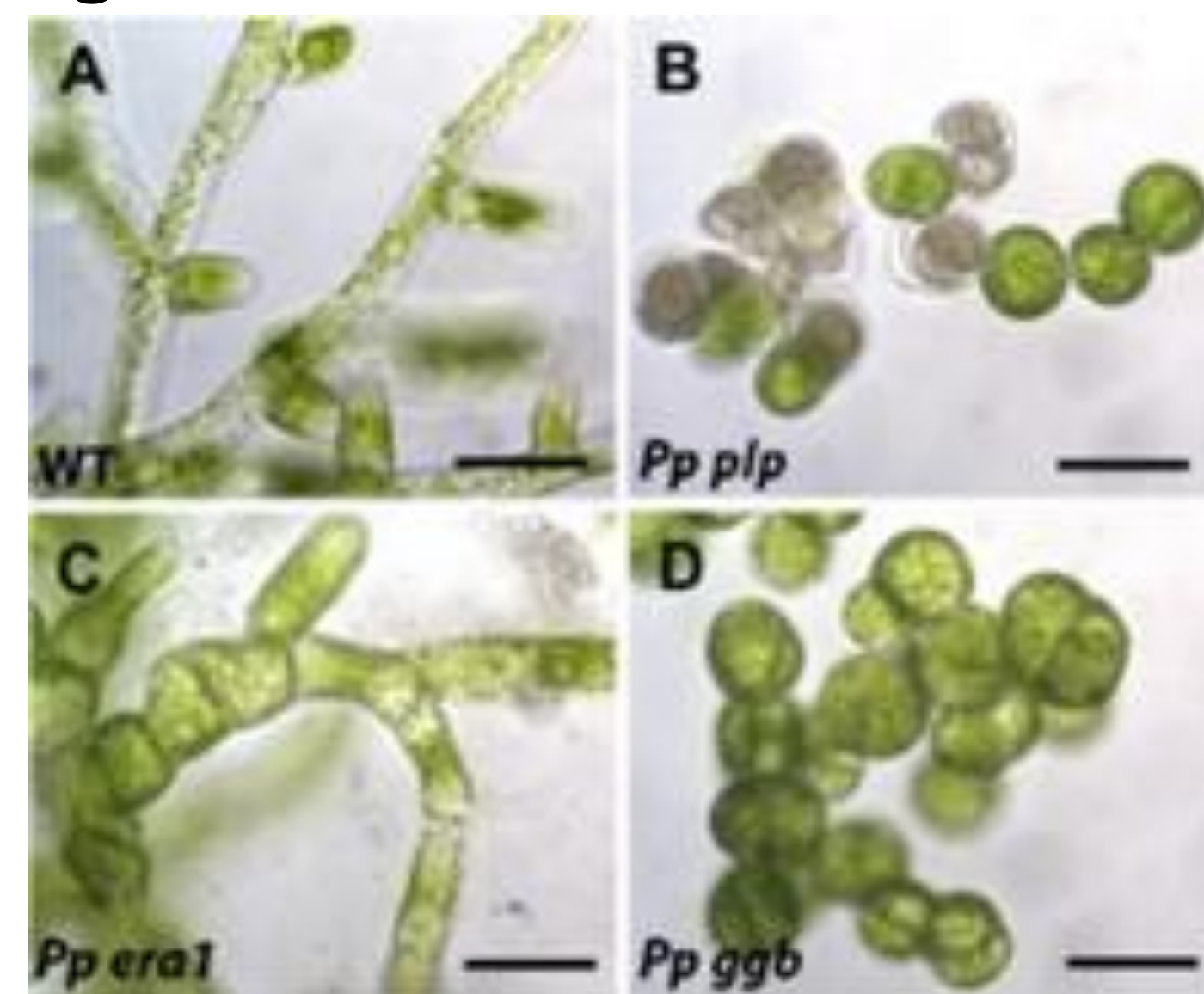
Introduction

Protein Prenylation is the addition of lipids to proteins and plays a vital role in the cell. It enhances protein-protein interactions and promotes protein-membrane targeting and association.

Protein Prenylation plays a key role in various plants' developmental processes. The key enzymes that are involved in prenylation are protein farnesyltransferase (PFT), protein geranylgeranyl-transferase-I (PGGT), and Rab geranylgeranyltransferase (Rab-GGT).

There has been prior research that examined the role of prenylation in plants. A study was done on Arabidopsis (*At*) and *Physcomitrella patens* (*Pp*) that analyzed the role of protein prenylation enzymes. A knockout of PGGT and PFT α and β subunit genes in *P. patens* resulted in phenotypic changes.

Figure 1



Wild Type (A) demonstrates elongation, filament formation, and adhesion which *Pp pfp* and *Pp ggb* (B and D) lack. *Pp era1* mutants show elongation but it is not as well as WT.

The knockout of *PpPPAL1* and *PpPPAL2* has resulted in the death of the cell. This suggests that these genes are necessary for moss growth and survival; however, their specific role is unknown. Previous studies have shown that prenylation plays a role in ABA response, polar cell elongation, and cell fate determination in Arabidopsis.

Further studies of the role of *PpPPAL1*, *PpPPAL2*, and prenylation in *P. patens* needed to be done. More specifically, whether these enzymes play a role in growth, hormone response, cell adhesion and determination, sugar response, and polar cell elongation.

Method

- To study the role of PPAL, a knockdown approach was used to reduce gene expression of *PpPPAL1* and *PpPPAL2* and characterize the resulting phenotype.
- Then, in-depth expression studies, metabolic analysis, and developmental studies of *P. patens* mutants was done.

Results

1. Reduction of gene expression of *PpPPAL1* and *PpPPAL2* was performed through gene knockdown:

- Artificial microRNA was created and used to target *PpPPAL1* and *PpPPAL2* genes
- *PpPPAL1* and *PpPPAL2* amiRNA transformants were grown in DMSO and β -Estradiol medium.

Figure 2. *PpPPAL1* Knockdown:



2. The most efficient moss transformants was used for RNA extraction and in-depth expression studies:

- DNA extraction from moss transformants and PCR to perform genotyping
- RNA extraction from moss transformants and perform RT-PCR to examine gene expression
- The most efficient moss transformants was selected via gel electrophoresis visualization.

3. Studying the developmental and metabolic process in *P. patens*:

- Cytoskeleton markers of knockdown plants was immunostained to examine the organization of cytoskeleton and cell division.
- RabA4b markers was transformed into knockdown plants to study vesicle transport and cell polarity.
- Certain metabolic process was analyzed to see if they are altered in the mutants.

Conclusion

- Study of gene expression of *PpPPAL1* and *PpPPAL2* knockdown lines will provide insights into the functional and conserved roles of *PpPPAL1* and *PpPPAL2* in *P. patens*.
- Preliminary results have shown that *PpPPAL1* and *PpPPAL2* knockdowns inhibit the growth and propagation of *P. patens*. It also resulted in less gametophore.
- More in-depth studies in *P. patens* will be done to test whether *PpPPAL1* and *PpPPAL2* and prenylation play a role in growth, hormone response, cell adhesion and determination, sugar responses, and polar cell elongation.

Significance

- Provide an understanding of how early plants adapt to land and the evolution of multicellularity.
- Can be used in agriculture to create plants that can tolerate drought.
- Provide uses in the field of algae biofuels by creating algae that can adhere to each other to lower the cost of harvesting.
- Provide an understanding of prenylation in humans, as defects in prenylation could lead to health problems.

Acknowledgement

I would like to thank the Undergraduate Mentored Research Award for providing the funds and Susana Perez Martinez, PhD student, and Dr. Mark P. Running for the opportunity to work on this project.