

LANDON SWARTZ

Graduate Student in Computer Science

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EXPERIENCE

Graduate Student

Computational Imaging and VisAnalysis Lab

Jan 2023 – Ongoing University of Missouri

- Novel research into computer vision for measurement of traits in life science and biomedical fields
- 3D aerial reconstruction

Undergraduate Researcher

David Mendoza-Cozatl Lab

Feb 2019 – Dec 2023 University of Missouri

- Automatic trait measurement of plants
- Embedded systems engineer for imaging robotics

PROJECTS

OPEN Leaf: Leaf Specific Phenotyping

National Science Foundation

March 2020 - Sept 2023 DMC Labs

- Designed a plant phenotyping pipeline that measured traits of shape and color of multiple plants at a leaf specific level to understand nutrient stressors

Segment-Anything Integration in SMART

National Science Foundation

Summer 2023 CIVA Labs

- Integrated the Segment-Anything Model (SAM) into a plant image analysis library (SMART) for improved trait measurement

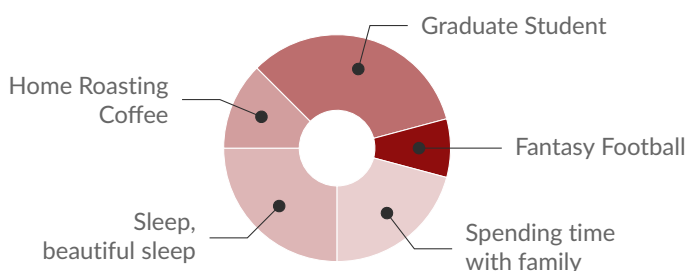
Leaf Tracking

National Science Foundation

Ongoing CIVA Labs

- Implemented tracking algorithm for leaves throughout time to perform dynamic trait measurement in the context of time

A DAY OF MY LIFE



MY LIFE PHILOSOPHY

"Trying to change the world for the better, one data set at a time."

MOST PROUD OF

OPEN Leaf Paper
being in the top 1% of papers on social media

Interdisciplinary
Apart of many cross-discipline teams for novel projects

TECHNICAL SKILLS

Machine Learning Statistical Analysis
Computer Vision Digital Image Processing
Precision Agriculture

C++ R Python PyTorch Open3D
Git OpenCV

SOFT SKILLS

Hard-working Eye for detail
Motivator & Leader Writer Personable

EDUCATION

Ph.D. in Computer Science

University of Missouri

Jan 2023 - Present

M.Sc. in Computer Science

University of Missouri

Jan 2023 - Present

- Expected date of award: Dec 2024

B.Sc. in Computer Engineering

University of Missouri

Aug 2018 - May 2022

- Minors in Mathematics and Biological Sciences
- Honors College Certificate

PUBLICATIONS

Journal Articles

- L. G. Swartz, S. Liu, D. Dahlquist, *et al.*, "Open leaf: An open-source cloud-based phenotyping system for tracking dynamic changes at leaf-specific resolution in arabidopsis," *The Plant Journal*, vol. n/a, no. n/a, DOI: <https://doi.org/10.1111/tpj.16449>. eprint: <https://onlinelibrary.wiley.com/doi/pdf/10.1111/tpj.16449>.

Conference Proceedings

- L. G. Swartz, K. Palaniappan, D. G. Mendoza-Cozatl, and S. Liu, "Segmenting arabidopsis leaves using segment-anything," in *52nd Annual Applied Imagery Pattern Recognition Workshop*, 2023.

PRESENTATIONS

The OPEN Series Design Philosophy, High Throughput Phenotyping in the Hands of Users

2023 North American Plant Phenotyping Network - 2 mins

 Feb 2023

 Danforth Plant Science Center, MO, USA

Open-Source Cloud Based Phenotyping Series (OPEN Leaf)

International Plant Sulfur Workshop - 15 mins

 July 2022

 Western University, Canada

Open-Source Cloud Based Phenotyping Series (OPEN Leaf and Root)

2022 North American Plant Phenotyping Network - 15 mins

 Feb 2022

 University of Georgia, GA, USA

POSTERS

Segmentation of Arabidopsis thaliana using Segment-Anything

2023 Artificial Intelligence and Pattern Recognition

 Sept 2023

 St. Louis University, MO, USA

OPEN Leaf: An open-source cloud-based plant phenotyping platform to track dynamic rosette changes at a leaf-specific resolution

2022 Interdisciplinary Plant Science Symposium

 May 2022

 University of Missouri, MO, USA

- Outstanding Poster Award

REFEREES

Distinguished Prof. Kannappan Palaniappan

 University of Missouri

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205 Naka Hall
Columbia, MO, USA

Prof. David Mendoza-Cozatl

 University of Missouri

 mendozad@missouri.edu

271f Bond Life Sciences Center
Columbia, MO, USA

COLLABORATIONS


OPEN Root

 2021 - Present

 DMC Labs

- Built an open-source high-throughput root phenotyper that captures initial Arabidopsis thaliana seedling root length and direction to study nutrient uptake for improved crop production

Synthetic Genetic Circuit Plants

 2022 - Present

 Dr. Jen Brophy, Stanford

- Conducted ionomics analysis of macro and micro nutrients on transgenic Arabidopsis thaliana lines created by Dr. Jen Brophy, assistant professor of bioengineering at Stanford, with synthetic genetic circuits to control root system architecture.

Yeast-II Hybrid

 2018 - Present

 DMC Labs

- Designed imaging platform for standardized image capture of yeast plates for transcription factor network analysis of entire genomes