Genomic and phenomic technologies for assessing root lesion nematode (Pratylenchus penetrans) resistance in red raspberry

Savannah Phipps^{1,2,3}, Inga Zasada^{1,2}, Michael Hardigan³, Jeff DeLong², Wendy Hoashi-Erhardt⁴, and Michael Dossett⁵

¹Department of Botany and Plant Pathology, Oregon State University; ²U.S. Department of Agricultural Research Service, Horticultural Crops Disease and Pest Management Research Unit; ¹Department of Botany and Plant Pathology, Oregon State University; ²U.S. Department of Agricultural Research Service, Horticultural Crops Disease and Pest Management Research Unit; ¹Department of Botany and Plant Pathology, Oregon State University; ²U.S. Department of Agricultural Research Service, Horticultural Crops Disease and Pest Management Research Unit; ³Department of Agricultural Research Service, Horticultural Research Service, ³U.S. Department of Agriculture–Agricultural Research Service, Horticultural Crops Production and Genetic Improvement Research Unit; ⁴Washington State University Puyallup Research and Extension Center; ⁵ British Columbia Berry Cultivar Development Inc.

Objectives

- Characterize root lesion nematode (RLN; Pratylenchus penetrans) resistance and tolerance phenotypes in a diverse population of 270 red raspberry breeding lines and cultivars and 10 mapping populations
- Determine the effectiveness of drone-collected spectral imaging data for assessing RLN damage
- Conduct genome-wide association study of RLN resistance using phenotypic and genotypic information
- Develop and evaluate genomic prediction models for improved selection of RLN resistance

Introduction

Management of root lesion nematode, *Pratylenchus* penetrans, (RLN; Fig. 1) is an important production goal for the profitable red raspberry industry in the unique climate of the Pacific Northwest (Figs. 2 & 3; Rudolph & DeVetter, 2015; Walters et al., 2017). Developing economical and sustainable control methods are a priority. Resistant cultivars are an effective control method; however, there is currently limited understanding of the genetics involved in host resistance. Furthermore, phenotypic evaluation of nematode resistance is an intensive and destructive process, which has hindered research into the genetics of RLN resistance. This project is currently evaluating newer technologies such as genomic prediction and



Figure 2: A red raspberry field in Northwestern Washington. Photo by Inga Zasada.

high-throughput imaging, which may accelerate screening and improve a breeder's ability to select leverages the combined

Photo by Inga Zasada. B) Pratylenchus spp. under a microscope from an for resistance. This project inoculated raspberry plant. Photo by Savannah Phipps. diversity of raspberry

Figure 1: A) Head region of

Pratylenchus penetrans.

germplasm from breeding programs at Washington State University (WSU), British Columbia Berry Cultivar Development Inc. (BCBCDI), the United States Department of Agriculture (USDA)/Oregon State University (OSU), and germplasm at the

National Clonal Germplasm Repository (NCGR) for assessment of RLN resistance.





Figure 3: Red raspberry 'Willamette' in a field with A) nonfumigated and B) fumigated plants with Pratylenchus penetrans present. Photos by Inga Zasada



Methods

Plant Materials

- 270 red raspberry genotypes from the WSU Small Fruits breeding program, BCBCDI, USDA/OSU breeding program, and NCGR
- 296 individual seedlings from 10 mapping populations derived from a susceptible by resistant cross made by the BCBCDI

Field Experimental Design

Figure 6 (above): A) Harvesting of

Phipps.

Baker.

aboveground biomass in year 1. B) Weighing

and recording of aboveground biomass in year

1. C) Weighing and recording of aboveground

Figure 7 (right): Examples of ground-based

and aerial imaging in action. A) Stitched red-

green-blue aerial orthomosaic of field trial from

May 5, 2023. Photo by Alexander Gregory. B)

MicaSense RedEdge-MX multispectral sensor.

Phipps imaging in the AgerPoint Capture app

biomass in year 2. Photos by Savannah

AgBOT mk 1 Quadcopter equipped with

Photo by Savannah Phipps. C) Savannah

with iPhone 14 Pro Max. Photo by Hannah

- Paired non-inoculated and RLN inoculated plants randomized across 3 replications (Figs. 4 & 5)
- Non-replicated seedlings from 10 mapping population planted alongside diversity panel (Fig. 4)



Figure 5: Examples of morphological diversity seen in field. A) 'Dorman Red' B) 'Jokgal' C) WSU 1478 D) 'Meeker'. Photos by Savannah

Phenotypic Data Collection

- Collected yearly
- Root sampling for nematode quantification
- Fresh weight aboveground biomass sampling (Fig. 6) for plant nematode stress response
- Spectral data collection via drone fly-overs and smart devices for plant nematode stress response (Fig. 7)

Genotypic Data Collection

Genotyping-by-Sequencing following the protocol outlined in Bushakra et al. (2015)

GWAS Analysis

FarmCPU in R with GAPIT (Lipka et al., 2012)

Genomic Prediction

Develop single and multi-year data models

Progress to Date

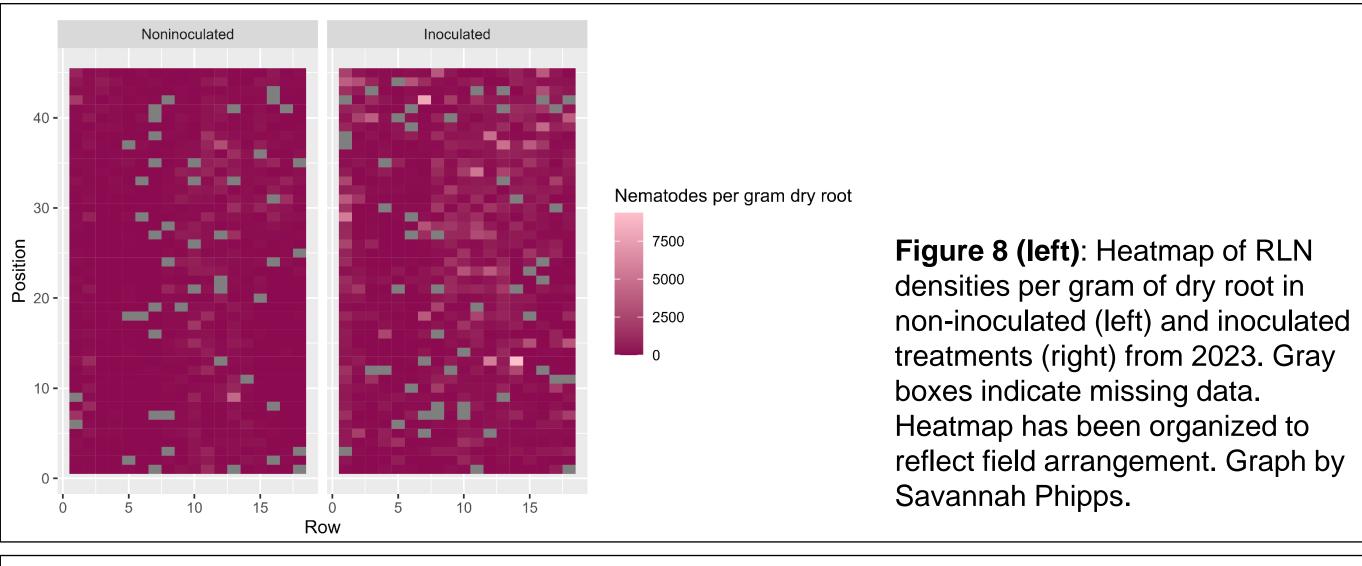
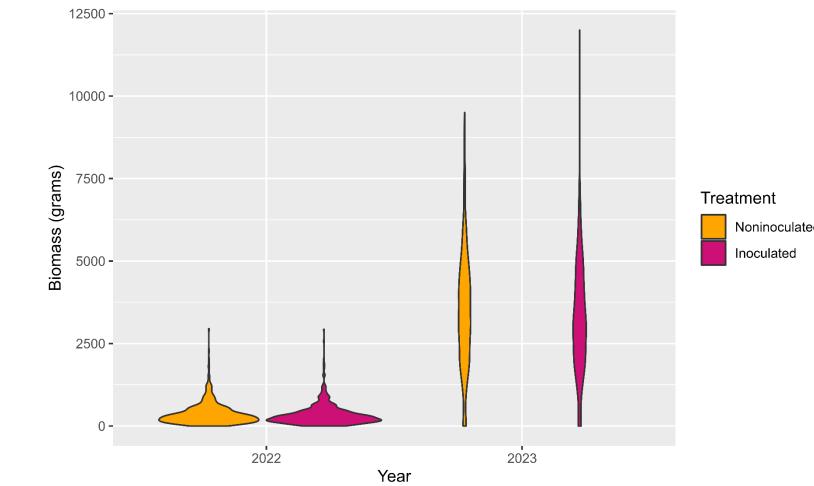


Figure 9 (right): Distribution of fresh weight aboveground biomass between RLN non-inoculated and inoculated treatments across trial years. Graph by Savannah Phipps.



- ❖ 1st year nematode quantification conducted (Fig. 8)
- ❖ 1st and 2nd year aboveground biomass measurements collected (Fig. 9)
- ❖ 1st year aerial and ground-based imaging conducted
- Initial data exploration and analysis ongoing
- Processing of image data ongoing

Future Work

- Final year of data collection
- Evaluate multi-year models for correlations between spectral data and biomass and nematode count data
- Conduct genome-wide association study with phenotypic and genomic data
- Evaluate genomic prediction models for multi-year data
- Publish results

Collaborations and Affiliations







Funding



Photo by Alexander Gregory

References

agerpoint

Bushakra, J. M., Bryant, D. W., Dossett, M., Vining, K. J., VanBuren, R., Gilmore, B. S., Lee, J., Mockler, T. D., Finn, C. E., & Bassil, N. V. (2015). A genetic linkage map of black raspberry (Rubus occidentalis) and the mapping of Ag_{4} conferring resistance to the aphid Amphorophora agathonica. Theoretical and Applied Genetics, 128, 1631-1646.

Lipka, A. E., Tian, F., Wang, Q., Peiffer, J., Li, M., Bradbury, P. J., Gore, M. A., Buckler, E. S., & Zhang, Z. (2012). GAPIT: genome association and prediction integrated tool. *Bioinformatics*, 28(18), 2397-2399. Rudolph, R. E. & DeVetter, L. W. (2015). Management strategies for Phytophthora rubi and Pratylenchus penetrans in floricane red raspberry (Rubus idaeus L.). Journal of the American Pomological Society, 69(3), 118-136. Walters, T. W., Bolda, M., & Zasada, I. A. (2017). Alternatives to current fumigation practices in western states raspberry. Plant Health Progress, 18, 104-111.