

Among population variation in root and shoot plasticity and plasticity of integration in *A. thaliana*



cousinsea@g.cofc.edu

Elsa A. Cousins and Courtney J. Murren
Department of Biology, College of Charleston, Charleston SC

murrenc@cofc.edu



Introduction

- Understanding of natural variation and stress tolerance in Arabidopsis can be applied across plant species³
- Root system architecture (RSA) and aboveground phenotypes are responsive to nutrient variation⁴
- Genetic variation in plasticity may be influenced by geographic origin^{1,6}
- Typically these ideas are addressed at the seedling stage without information about ecological soil history

Study Populations

Do geographically close yet distinct populations respond to the same treatments in similar ways in both roots and shoots?



Methods

Seed stock selected and planted in December 2014

Greenhouse Nutrient Treatments: **Promix**, **Hoagland's**, **Low Phosphorous**, **Low Nitrogen**
Agar experiment: complete nutrient medium

Greenhouse experiment:

- bolting date, rosette diameter, fruit number, height
- harvested when basal fruits were mature
- roots cleaned and scanned with Win-RHIZO, analyzed for average diameter and total length

Agar experiment:

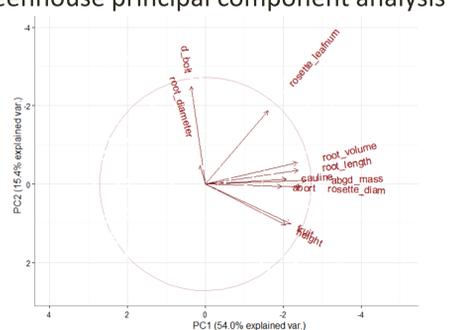
- 14 day old seedlings
- roots and shoots massed on micro-balance
- trays scanned with Win-RHIZO, analyzed for average diameter and total length

Trait integration varies by population and environment

Greenhouse principal component analysis

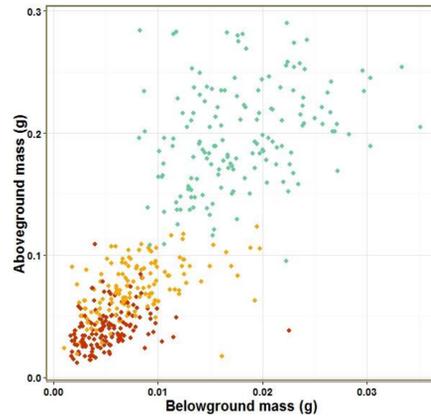
PCA for plants grown in sand treatments:
Populations and treatment were highly significantly different for both PC1 and PC2

- PC1:
- population cov. est. 0.24 (0.10) p<0.01
 - treatment F=1450.3 p<0.0001
- PC2:
- population cov. est. 0.67(0.26) p<0.0001
 - treatment F=10.81 P<0.0001



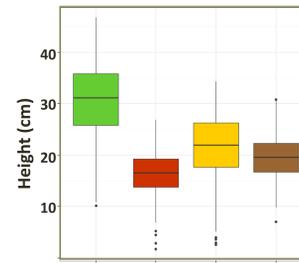
Plasticity to nutrient environments in trait means and trait relationships

Aboveground and belowground mass and relationships vary by treatment



Low Phosphorous: spearman 0.52, p<0.0001
Low Nitrogen: spearman 0.43, p<0.0001
Hoagland's: spearman 0.35, p<0.0001

Plant height varies by treatment

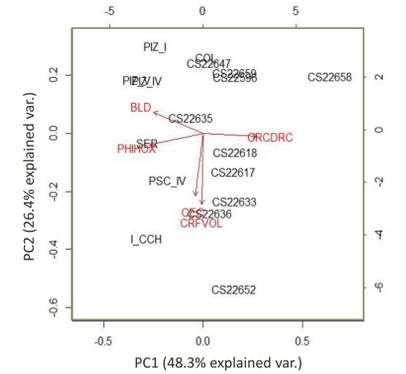


Treatments
Color Key:
Hoagland's
Low Phosphorous
Low Nitrogen
Promix

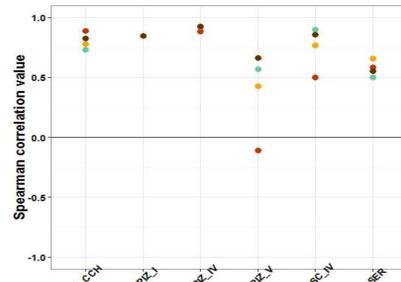
Soil quality of origins

Edaphic factors are predictive of integration and plasticity in seedling and adult plants

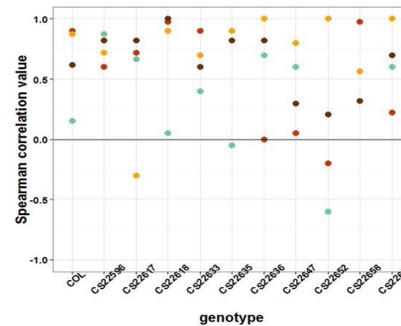
Soil principal component analysis



Genetic and environmental variation in aboveground traits: variation in integration and plasticity in lines with differing soil histories

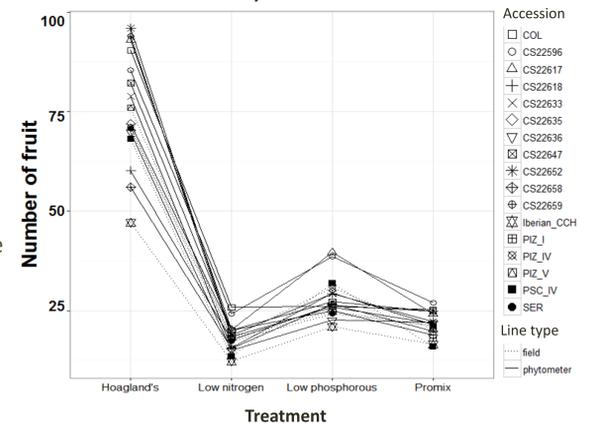


Fruit and height correlations



- Traits do not vary independently
- Interrelated functions have the potential to influence evolution^{2,5}
- Populations from similar soils have similar correlations

Fruit number varies by accession and treatment

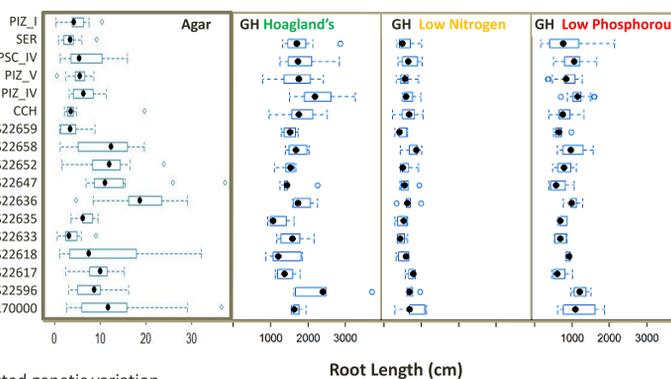


We found plasticity for fitness in total reproductive output (F_{3,45}=294.5, p<0.0001)

- Significant variation in reproductive output:
- accession (covariate estimate 61.8±35, p<0.005)
 - accession by treatment (covariate estimate 58.2±24.2, p<0.01)

Seedling root systems do not directly correlate with adult phenotypic variation and plasticity

Agar and greenhouse root length varies by accession



We detected genetic variation across accessions (F_{16,207} = 4.98 p<0.0001) in seedlings

We detected genetic variation across accessions (p<0.001), variation across treatments (F_{2,30} = 227.6 p<0.001), and genetic variation for plasticity (p<0.0008) in mature plants

The effect of landscape level environmental variation is observed in population differentiation

Developmental stages need to be considered when evaluating root traits as patterns in RSA can vary by:

- life stage
- environment
- genotype

Acknowledgments

Thanks to Rebecca Balazs and Claire Kohler and all of the Murren and Rutter labs and the unPAK team for their help, especially Bravada Hill, Quinten Meadors, and Gorka Sancho as well as all others involved in the curation of the seed stock of the Iberian populations. Follow the unPAK project at arabidopsisunpak.org

Literature cited

1. Agren, J., Oakley, C., McKay, J., Lovell, J., & Schemske, D. (2013). Genetic mapping of adaptation reveals fitness tradeoffs in Arabidopsis thaliana. *Proceedings of the National Academy of Sciences*, 21077-21082
2. Armbruster, W.S., Pélabon, C., Bolstad G.H., Hansen T.F. (2014). Integrated phenotypes: understanding trait covariation in plants and animals. *Philosophical Transactions of the Royal Society B* 369: 20130245.
3. Den Herder, G., Van Isterdael, G., Beekman, T., De Smet, I. (2010). The roots of a new green revolution. *Trends Plant Sci* 15: 600-607.
4. Julkowska, M., Hoeksloot, H., Mol, S., Féron, R., de Boer, G., Haring, M.A. & Teuberink, C. (2014). Capturing Arabidopsis Root Architecture Dynamics with ROOT-FIT Reveals Diversity in Responses to Salinity. *Plant Physiology*, 166, 1387-1402.
5. Murren, C.J. (2012). The Integrated Phenotype: Integrative and Comparative Biology 52: 64-76.
6. Weigel, D. (2012). Natural Variation in Arabidopsis: From Molecular Genetics to Ecological Genomics. *Plant Physiology*, 2-22.

Funding

IOS1146977 to Murren
IOS1052262 and IOS1355106 to Rutter, Strand, Murren

