Taming Variation:

Wild-types, Ecotypes and Phytometers of Arabidopsis thaliana and the UNPAK Project

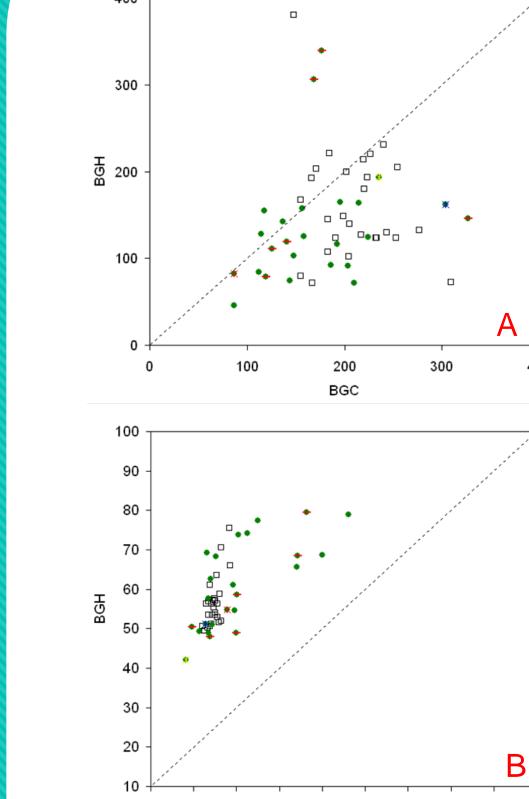
HS Callahan¹; CC Kohler^{1,2}; CJ Murren²; MT Rutter,² ; A Strand²; M Wolyniak³; and a large team of undergraduates ¹Barnard College, ²College of Charleston, ³Hampden-Sydney College

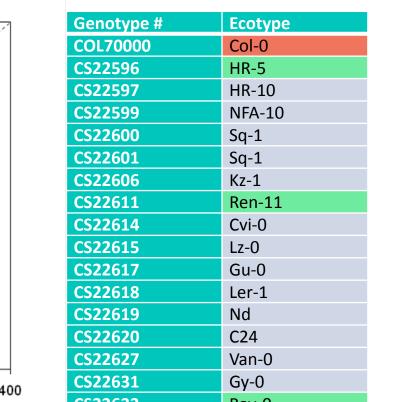
UNPAK : Quick Overview

Our collaborative network is phenotypically characterizing a large fraction of the extensive SALK knockout mutants library in Arabidopsis thaliana, emphasizing fitness-related traits and evolutionary questions, asking

- How often and which mutations result in no change, an increase, or a decrease in fitness-related traits? This information is a significant gap in our understanding of gene and genome evolution.
- Are fitness-related phenotypes associate with

From Ecotype to Phytometer: Screening of Candidate Accessions



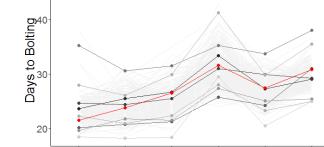


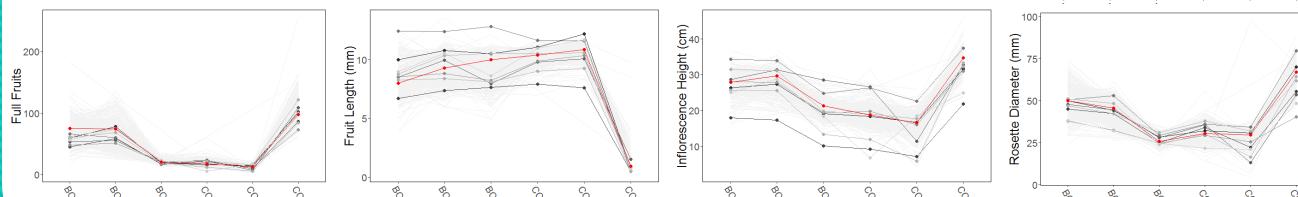
Beta Testing: Phytometer Performance Across Blocks in a Large-Scale Mutant Screen

An early UNPAK experiment was blocked spatially (BC, Barnard; CC, College of Charleston) and temporally, and phenotyped 362 SALK knockouts (gray lines), 10 phytometers (lines with points) to wild type and variable phytometers. and Col-0 wild-type (red).

Experiment Name

The trait-specific phenotypic plasticity elicited by unplanned environmental variation. can be addressed statistically and further contextualized by comparing





variation in several gene attributes? We can consider ontology, gene family size, many other attributes.

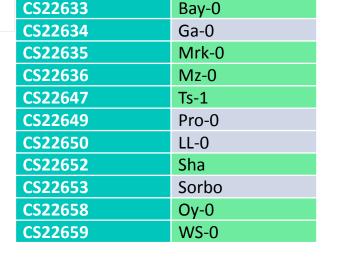
- Can we maximize undergraduate engagement in \bullet authentic genomics research? Beyond involvement in plant care and experiment implementation, they also design experiments, manage data, conceive and execute analyses, collaborate and communicate.
- Novel gene discovery or elucidation of gene function is not a central goal, but may be a by-product of UNPAK's efforts.

A Few Key Terms and Concepts

Genotype $\leftarrow \rightarrow$ Phenotype

"code"	\leftrightarrow	"function" (gain or loss)
"Normal" allele	$\leftarrow \rightarrow$	Wild-type function or trait
Mutation in allele	\leftrightarrow	Mutant phenotype
Natural polymorphic allele(s)	$\leftarrow \rightarrow$	Natural phenotypic variant(s)
accessions" – sequences vary	\leftrightarrow	"ecotypes" – traits vary

Knockout mutant library DNA library $\leftarrow \rightarrow$ Gene pool (evolution) $\leftarrow \rightarrow$ Population (functional ecology)



We screened 27 candidate phytometer lines (green dots) selected from an initial 97 lines. Pilot experiments included 27 SALK KO (open squares).

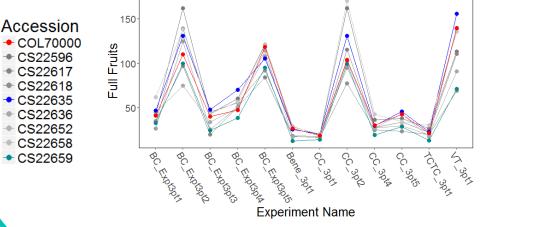
Phenotyping UNPAK traits was completed in growth chamber (BGC) and greenhouse (BGH) experiments, scoring means for fruit length, full and aborted fruit counts (and total fruits, A), inflorescence height and branching, rosette diameter, days to bolting (B) and germination. We selected the final panel of 10 phytometers (green in table), favoring ecotypes that bracketed variation in one or more traits expressed in SALK knockout library lines, created in a Col-0 wild-type background.

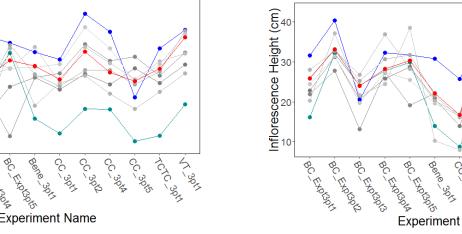


Plasticity was elicited by soil treatments, and was similar in magnitude to that elicited by unplanned variation in previous studies that included campusto-campus and temporal blocks

6 soil treatments phytometer + wt 62 SALK KO lines N = 2,595 pots

Here Come the Phytometers: Performance Across UNPAK Campuses Phytometers are helpful with Plots documenting mean phytometer performance any inconsistencies in through time, before and phenotyping, important given after expansion of UNPAK to UNPAK's reliance on multiple Pls and undergraduate interns. many more campuses.





Take-homes: On Mutants, Wild-types and Phytometers

- Concept of "a gene" in isolation is useless and meaningless
- Gene-trait-environment combinations are key



Genome $\leftarrow \rightarrow$ Phenome

Is a mutant deviant, compared to natural genotypes? How plastic are mutants and wild-types to change in the micro-How plastic are environment mutants or wild-types to change in macro-How relevant is environments? phenotypic variation and

plasticity detected in lab settings or mutants?

• Soil treatments revealed some trait deviation in KO lines, but not more often than prior experiments

Key References

696.

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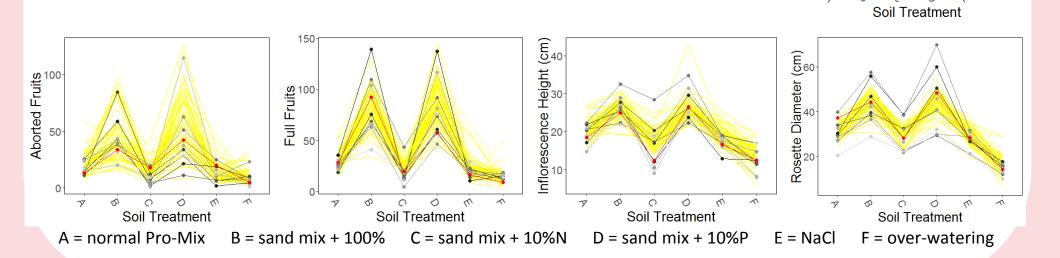
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Microspatial heterogeneity of the edaphic

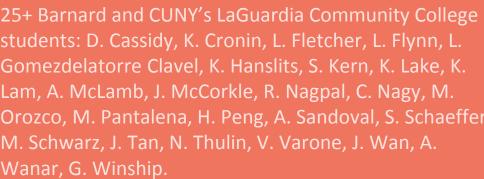
environment. Journal of Ecology 79: 687-

of leaf growth and molecular phenotypes.



- Environmental variation can be neither controlled nor eliminated
- UNPAK's phytometers are not universal. They match our core traits and environments. A new set may be needed for other traits or environmental conditions.
- Compare to agri-business "bundles" –seed strain and accompanying fertilizer/herbicide/pesticide must match local climate and soil
- Key concept: distinguish "signal" from "noise" in data about phenotypes
- Phytometers, like any other well-characterized and useful biodiversity, are similar to intellectual property such as software

UNPAK: Plant genomics on the shoulders of undergraduates !!











Lessons from 200,000 Arabidopsis phenotypes: the power of UNPAK

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Come to Matt Rutter's talk, Tues @ 9:00 AM, **Session 6B**





