

Using Enviromics to Identify More Adapted Angus Cattle



Troy Rowan
University of Tennessee
BIF Adaptation & Efficiency Breakout
June 11, 2024

Animal Performance

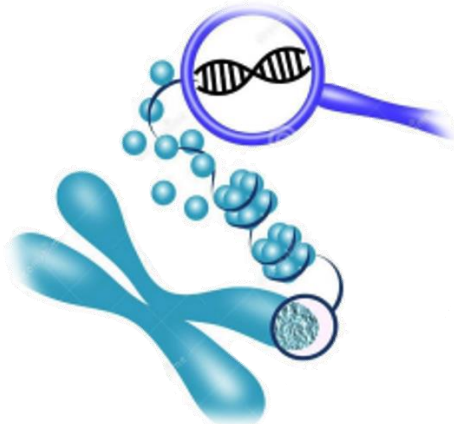
Phenotype ← Genetics & Environment

Production
Reproduction
Disease resistance
Product quality



Phenomics

Breeds
Lines
Individual variation



Genomics

Climate
Diet
Management
etc.



Enviroomics

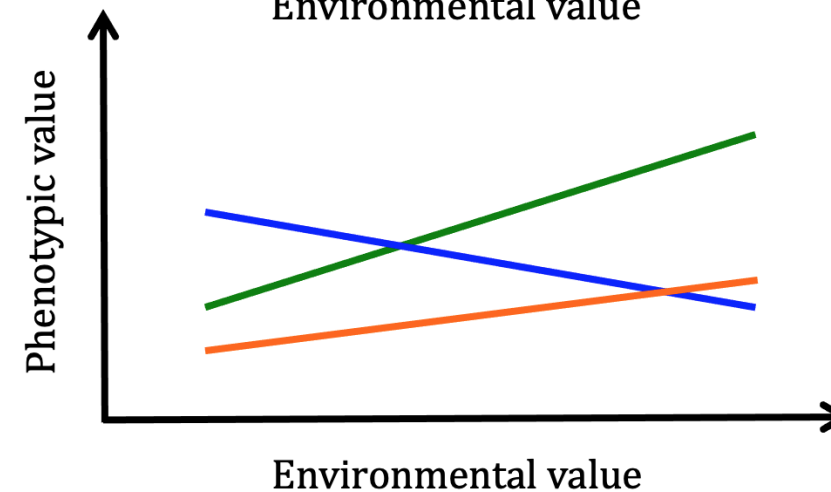
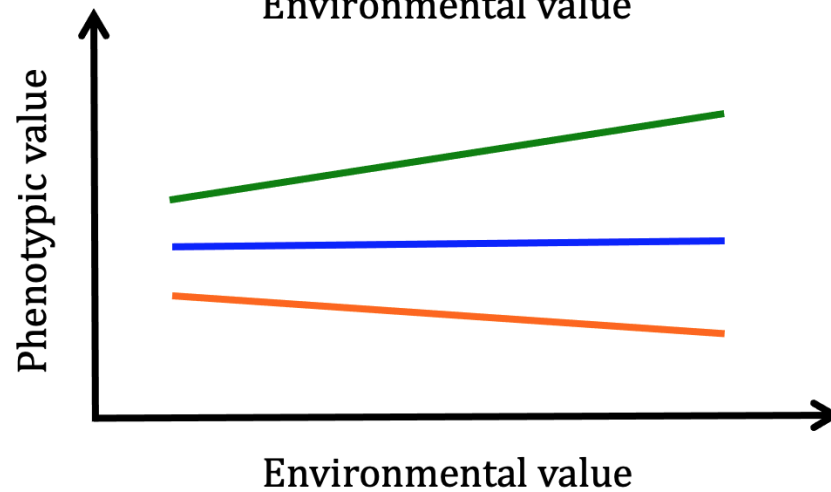
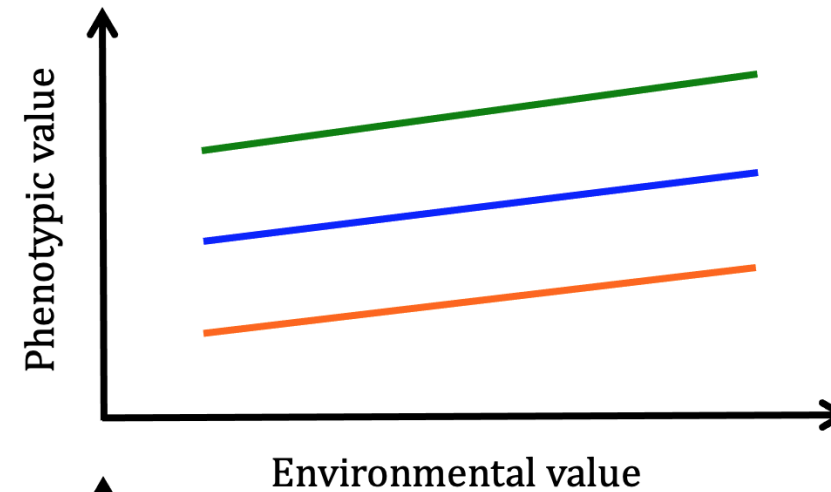
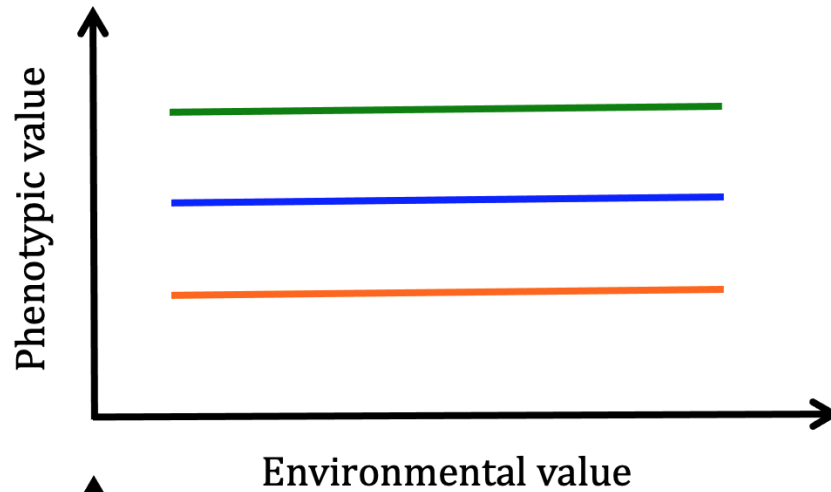
Optimization of Genetics, Management and Environment Combination



Genotype x Environment



Genotype x Environment (Plasticity)



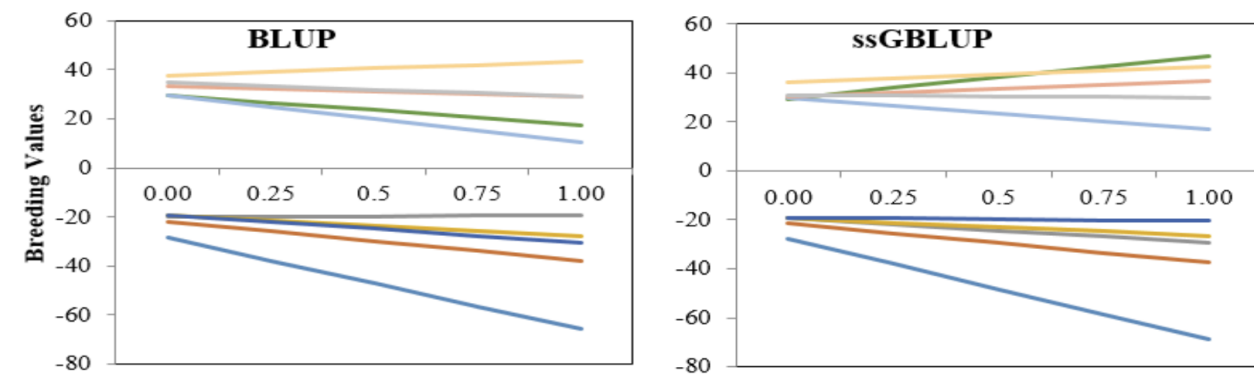


Figure 1. Estimated breeding values from BLUP and ssGBLUP for the five best and worst bulls for yearling weight in Nellore cattle over the environment gradients. Adapted from Oliveira *et al.* (2018).

What we have done:

- Single trait/env reaction norms
- Heat stress
- Regional genetics correlations
- Adaptive traits

What we haven't done:

- Dense environmental characterization
- Disentangled GxM from GxE
- Multi-trait/multi-variable work

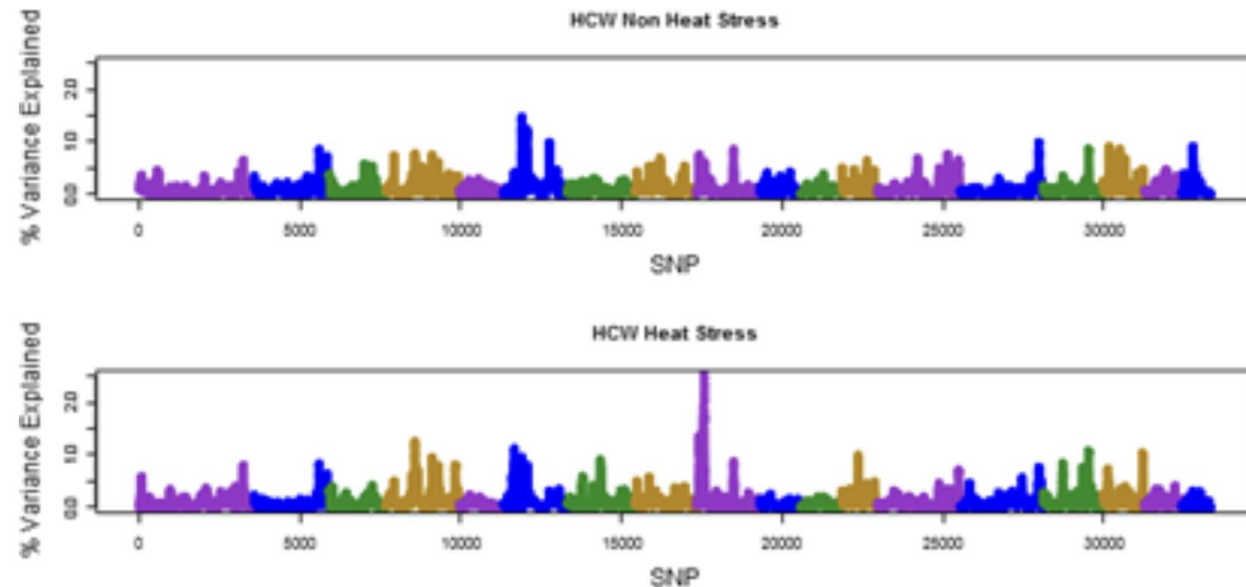


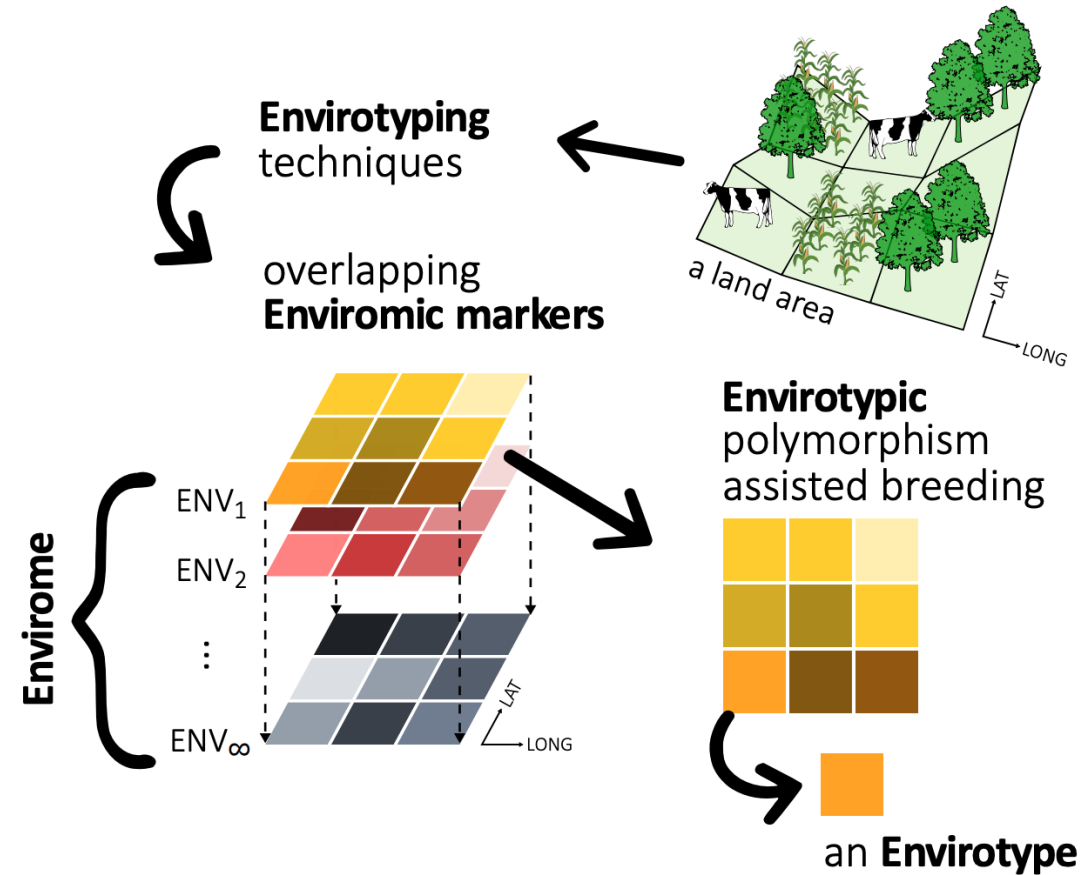
Figure 2. Manhattan plots for hot carcass weight (HCW) under no heat stress and heat stress. Chromosomes are indicated with different colors.

G x E Analysis

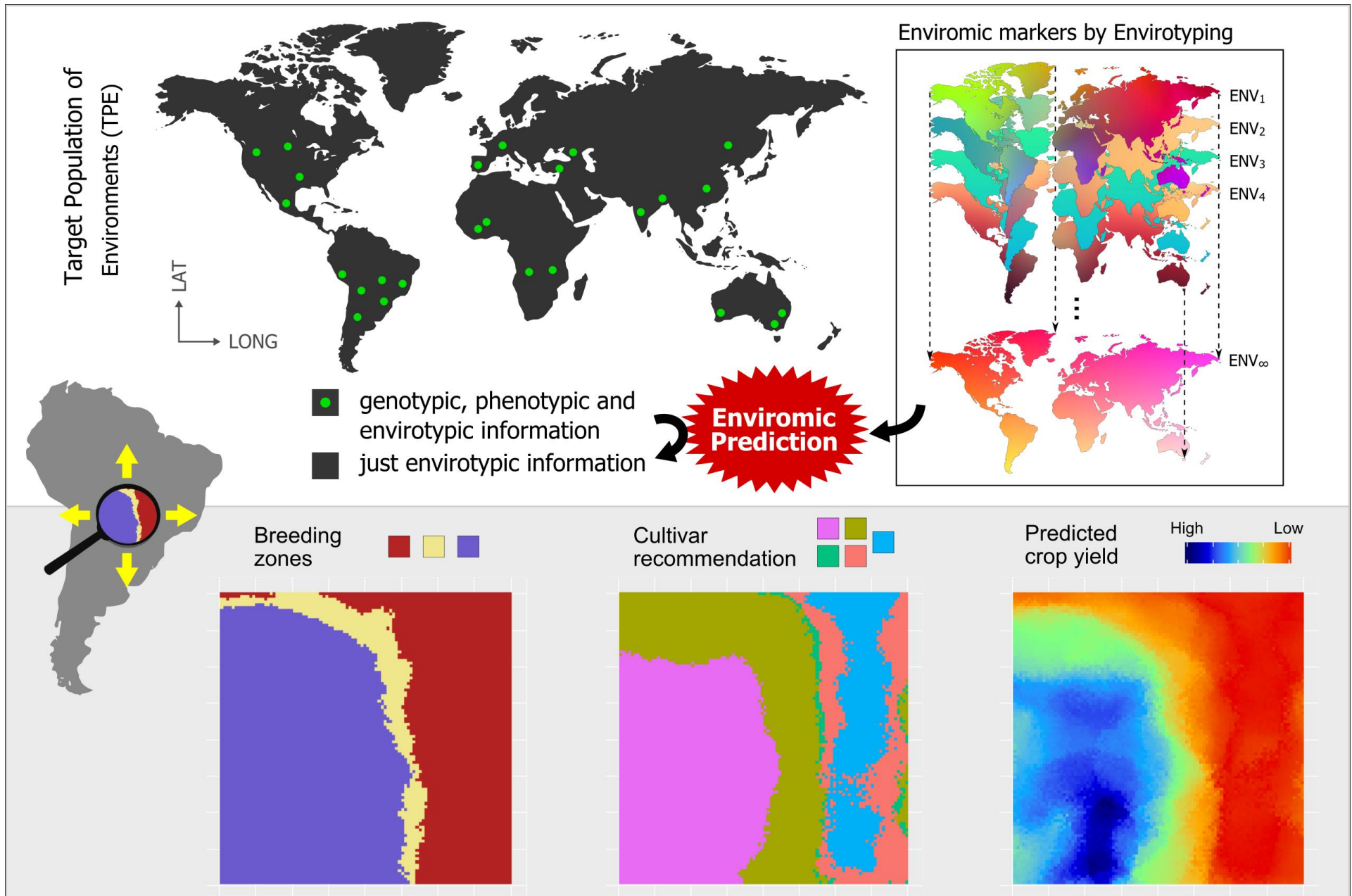
- Visualization Tools and Modeling Approaches
 - Finlay–Wilkinson, AMMI, GGE biplot
 - Interaction term ($P = G + E + G \times E + e$)
 - Multi-trait models
 - Reaction norms



Enviromics in Breeding



Resende, R. T., Piepho, H.-P., Rosa, G. J. M., Silva-Junior, O. B., Silva, F. F., Resende, M. D. V and Grattapaglia, D. Enviromics in breeding: applications and perspectives on envirotypic-assisted selection. *Theoretical and Applied Genetics* 134: 95–112, 2021.





Integrating Enviromics, Genomics, and Machine Learning for Precision Breeding of Resilient Beef Cattle

(USDA AFRI 2023-68014-39816)



What does a zip code miss?



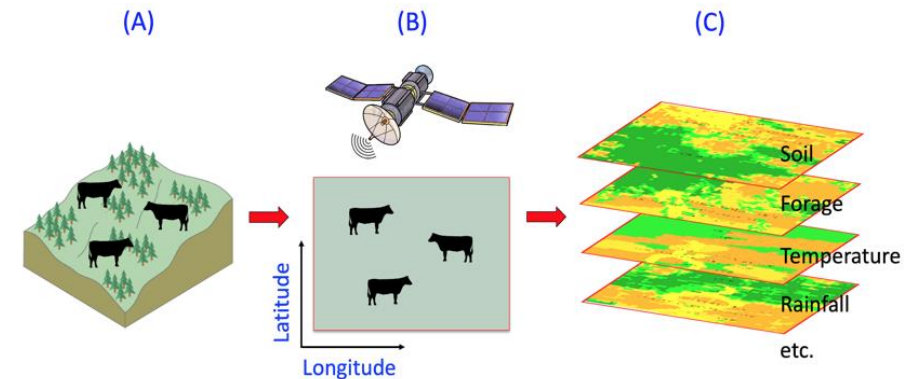


Integrating Enviromics, Genomics, and Machine Learning for Precision Breeding of Resilient Beef Cattle

(USDA AFRI 2023-68014-39816)

Research-Extension Integrated Objectives:

1. Generation of data lake and data processing pipelines for comprehensive environmental characterization of US beef cattle production systems.
2. Comprehensive evaluation of genotype-by-environment interactions and future performance through an enviromics approach.
3. Definition of novel indicators of animal resilience through in-depth phenotyping of genetically divergent animals.



WISCONSIN
UNIVERSITY OF WISCONSIN-MADISON

Rosa, Huang



UNIVERSITY OF
GEORGIA

Lourenco



THE UNIVERSITY OF
TENNESSEE
KNOXVILLE

Rowan



PURDUE
UNIVERSITY

Brito, Johnson



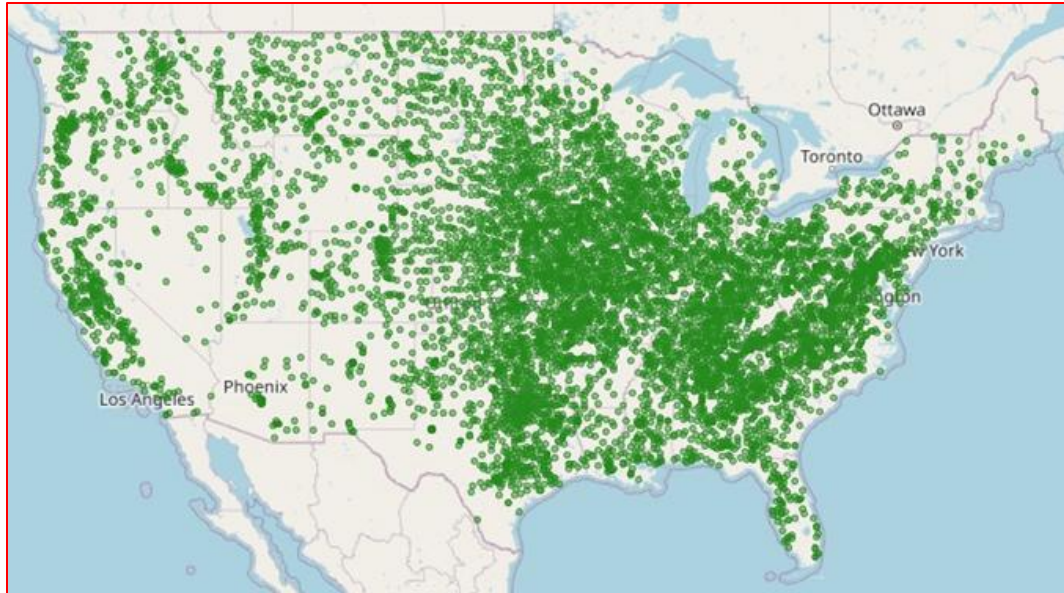
MICHIGAN STATE
UNIVERSITY

Gondro, Valle de Souza



Retallick

Overall Project Structure



Farm management



Soil, climate, and weather information

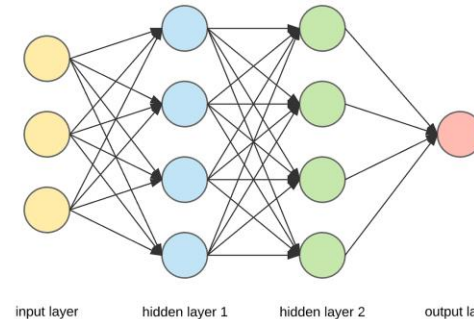


Forage density and quality

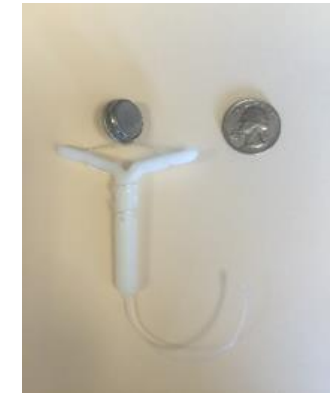


- 9K+ herds (1970-2020)
- 9.3M+ birth weights
- 9.8M+ weaning weights
- 4.8M+ post-weaning weight gain
- 126K+ heifer pregnancy
- full pedigree information
- 1M+ genotypic information

Data analytics



Extension and outreach

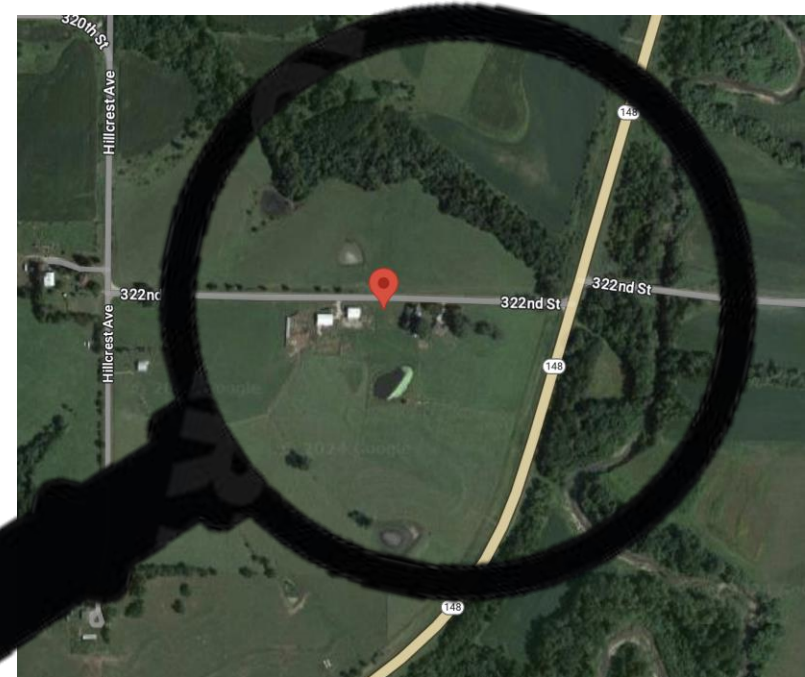


Novel indicators of animal resilience

How do you understand management? Ask about it!

Developing a survey to refine intensity of management:

- Forage and Grazing Management
- Health
- Reproduction
- Business Decisions
- Technology adaptation
- Genetic Decision-Making
- Demographics



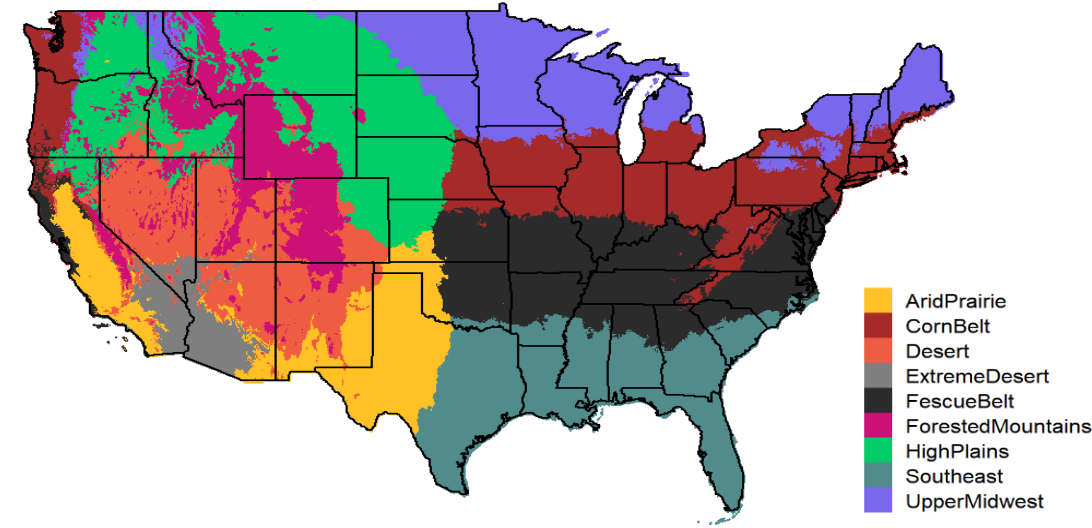
How will we model GxE(xM)?

Regional GxE(xM) genetic correlations

Multi-Dimensional Reaction Norms

Geographically Weighted Regression (GWR)

Ensemble of Machine Learning approaches



Thanks for Listening!

Angus Producers: Stay tuned for communication about survey rolling out in July/August!

Reach out with questions and ideas!

trowan@utk.edu

(712) 427-0096

Guilherme Rosa (Project PI)

grosa@wisc.edu

