

# Genotype by environment interaction in the 2014 Inbred Study

Celeste Falcon

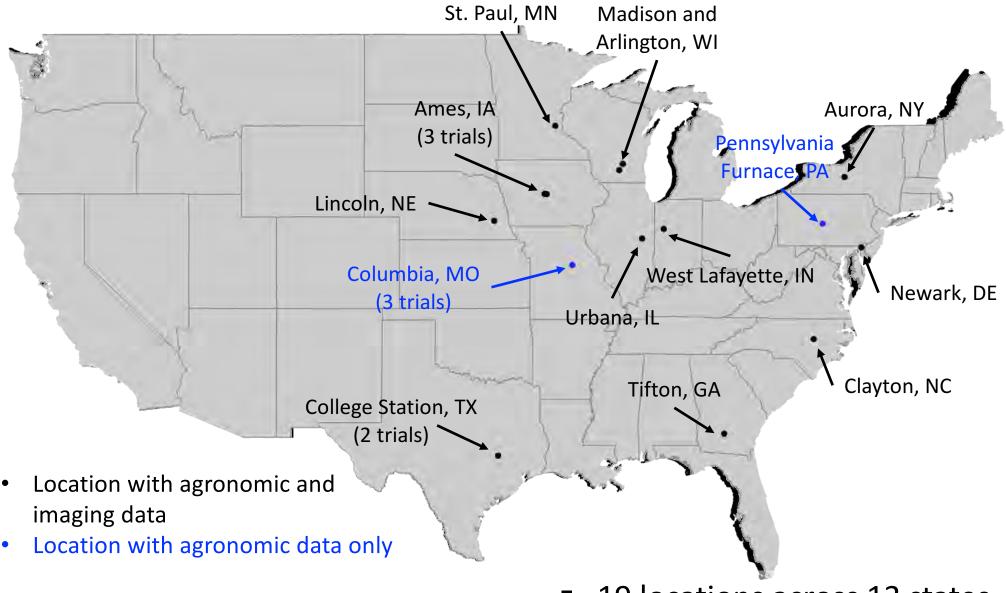
ASTA meeting
December 8, 2016

www.Genomes2Fields.org

## 2014 GxE inbred study—Questions to explore

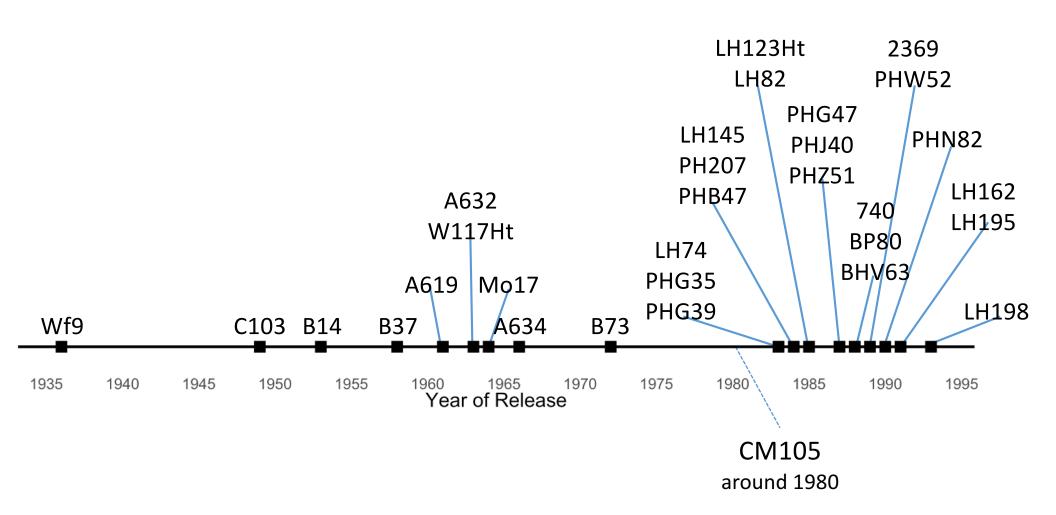
- What correlations do we observe among agronomic and yield component traits?
- ♦ How do trait means change with respect to inbreds' year of release?
- ♦ For which traits does GxE interaction explain a significant portion of variation observed?
- ♦ What is the distribution of GxE among traits?
- ♦ Which inbreds show the most stability?
- ♦ How does GxE change with respect to inbreds' year of release?

#### 2014 Inbred trials



- 19 locations across 13 states
- 2 replicates per environment

## Population: 31 inbred lines selected to represent range of locations and release date

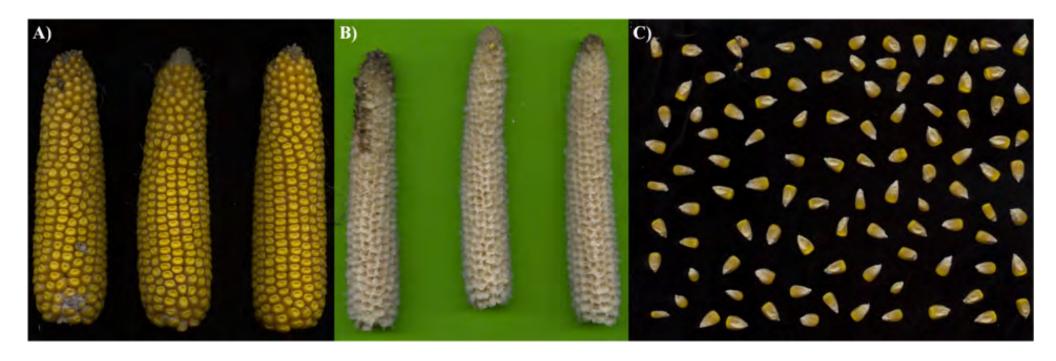


#### Phenotypic data

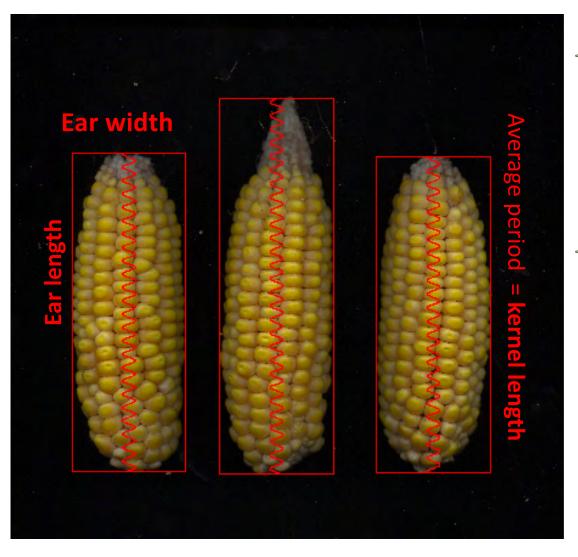
- **♦** Agronomic traits
  - ♦ Anthesis date (DAP)
  - **♦**Silking date (DAP)
  - **♦**Plant height

- **♦**Yield component traits
  - **♦**Ear length
  - **♦**Ear width
  - ♦ Kernels per row
  - **♦**Kernel row number
  - **♦**Kernel weight

## Yield Components: Imaging Output



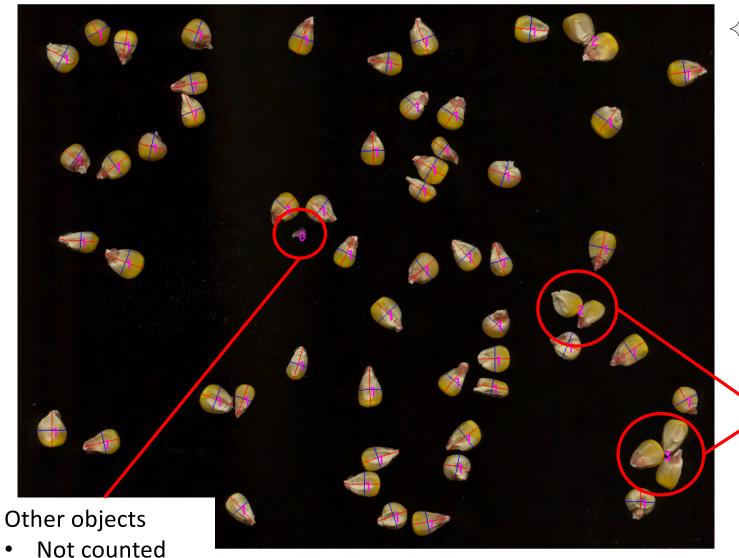
#### **Imaging Output: Ears**



- **◇Kernel row number**: counted manually

#### Imaging Output: Kernels

60



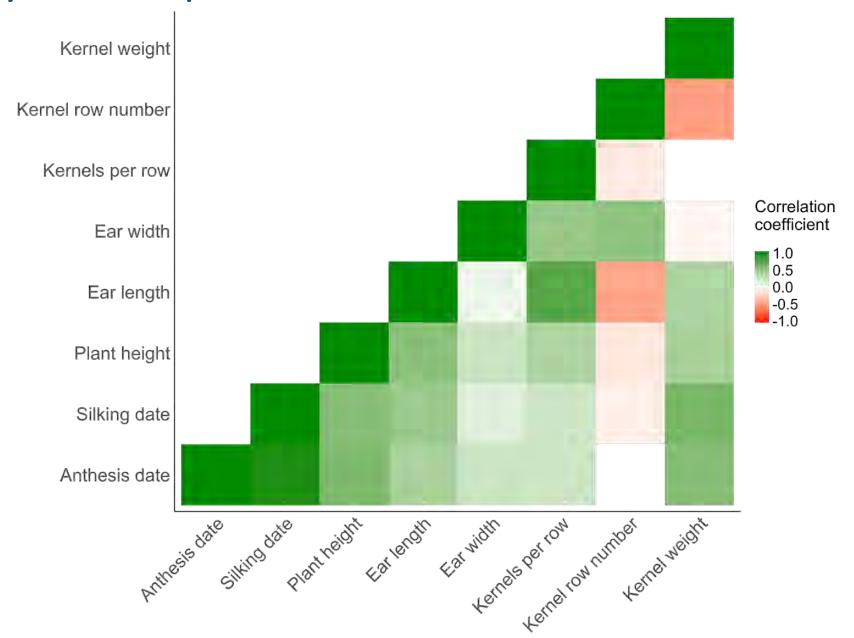
**Kernel weight** = Cup weight / kernel count

Grouped kernels:

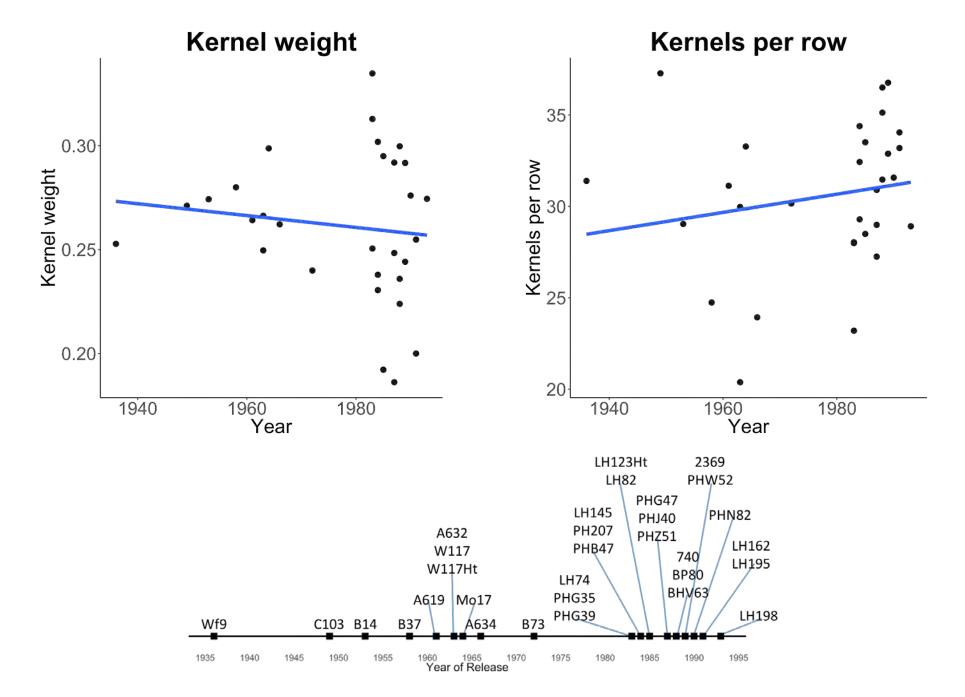
- **♦** Counted accurately
- ♦ No measurements recorded

- Not measured

## What correlations do we observe among agronomic and yield component traits?



#### How do trait means change w.r.t. year of release?



#### Quantifying GxE interaction

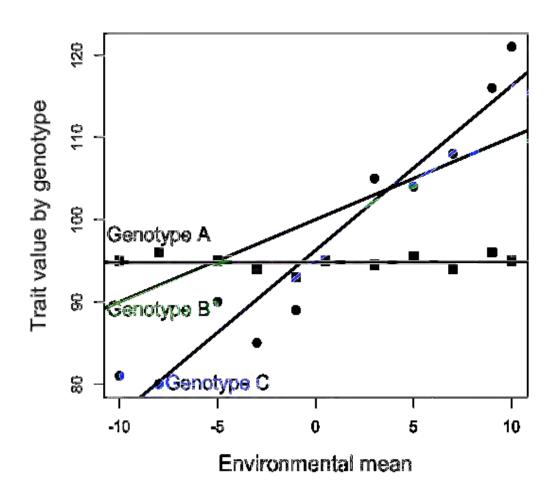
#### **♦Slope**

- $\Rightarrow$  = 0: Type I stability

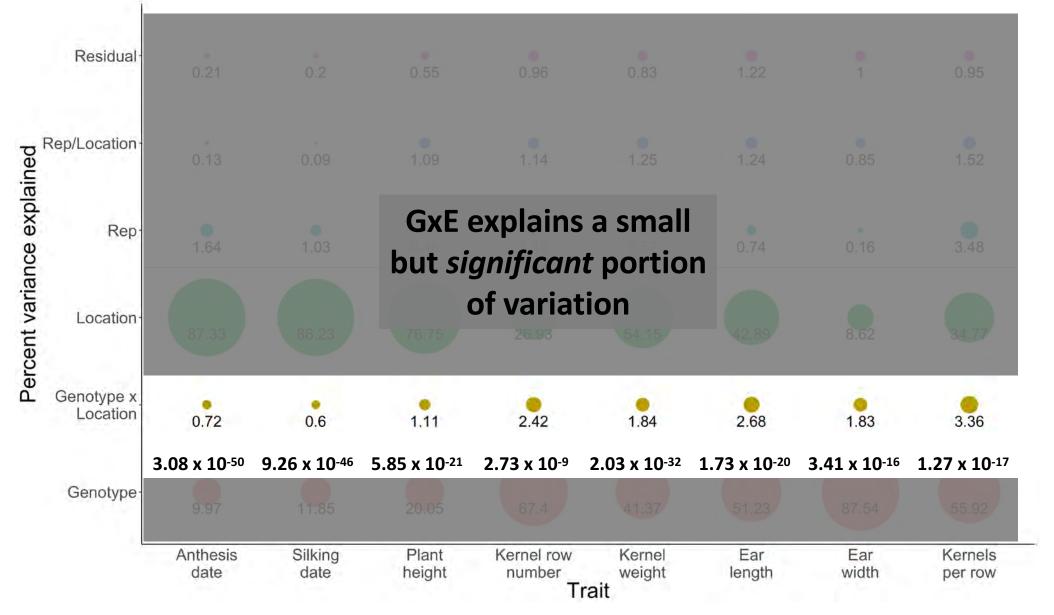
  - ♦ Ex) Genotype A
- ♦= 1: Type II stability

  - ♦ Ex) Genotype B
- ♦ Mean square error
  - ♦ Low MSE: Type III stability
    - ◆ predictable performance across environments

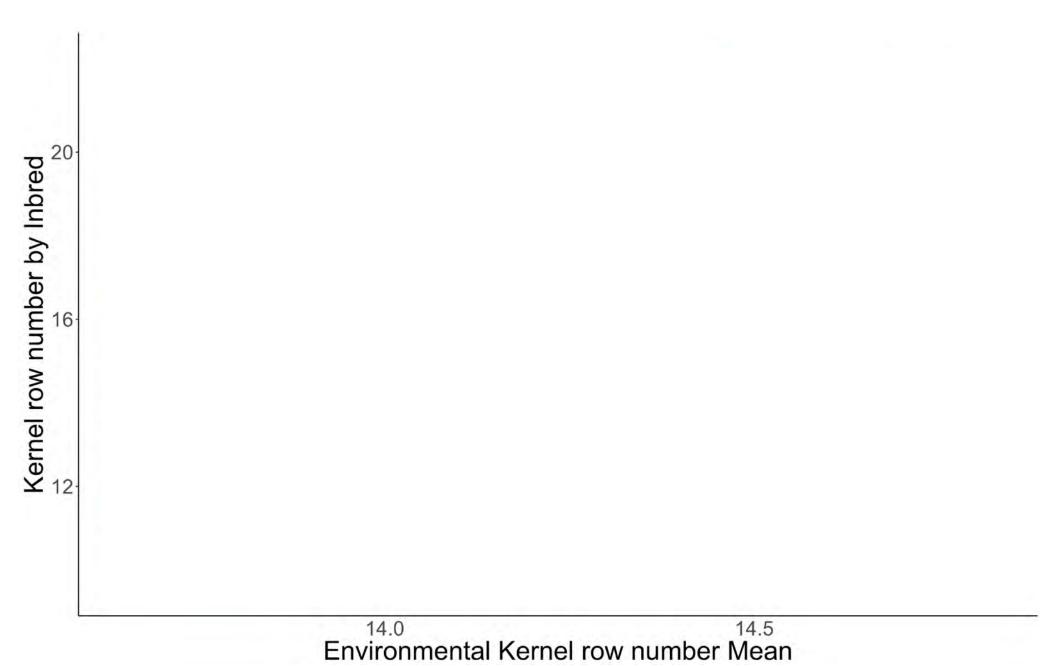
Standardized MSE = MSE / mean trait value<sup>2</sup>



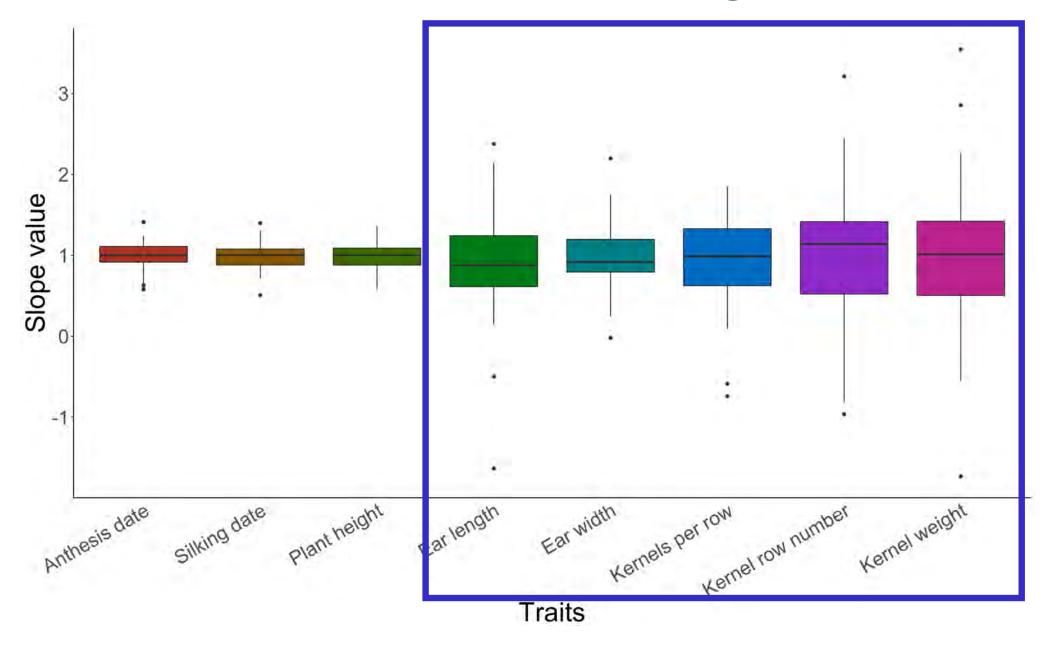
## For which traits does GxE interaction explain a significant portion of variation observed?



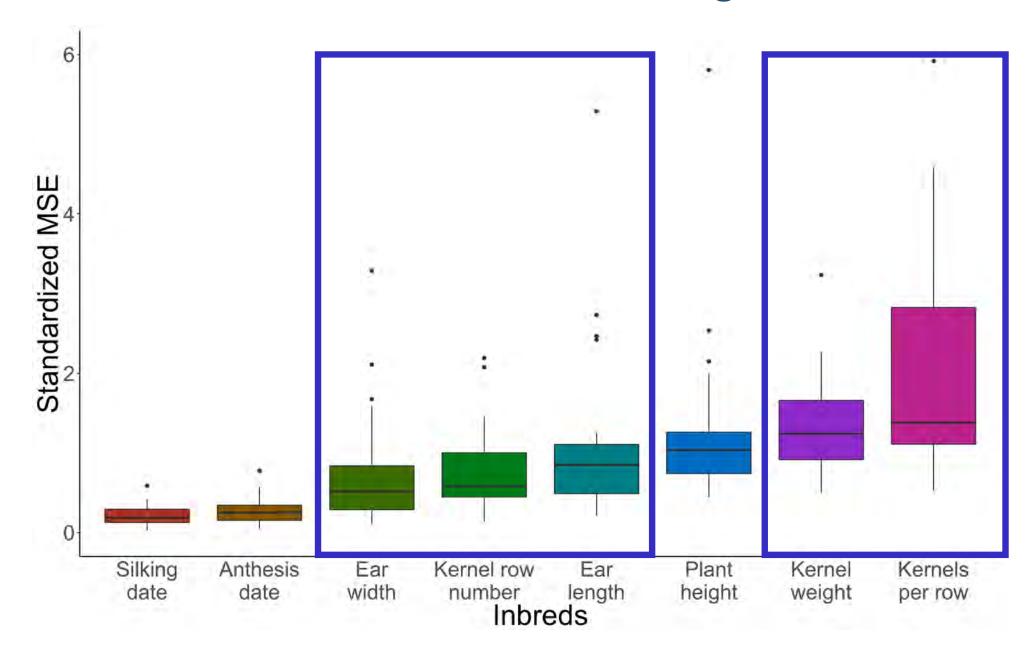
## **GxE Regression Plot**

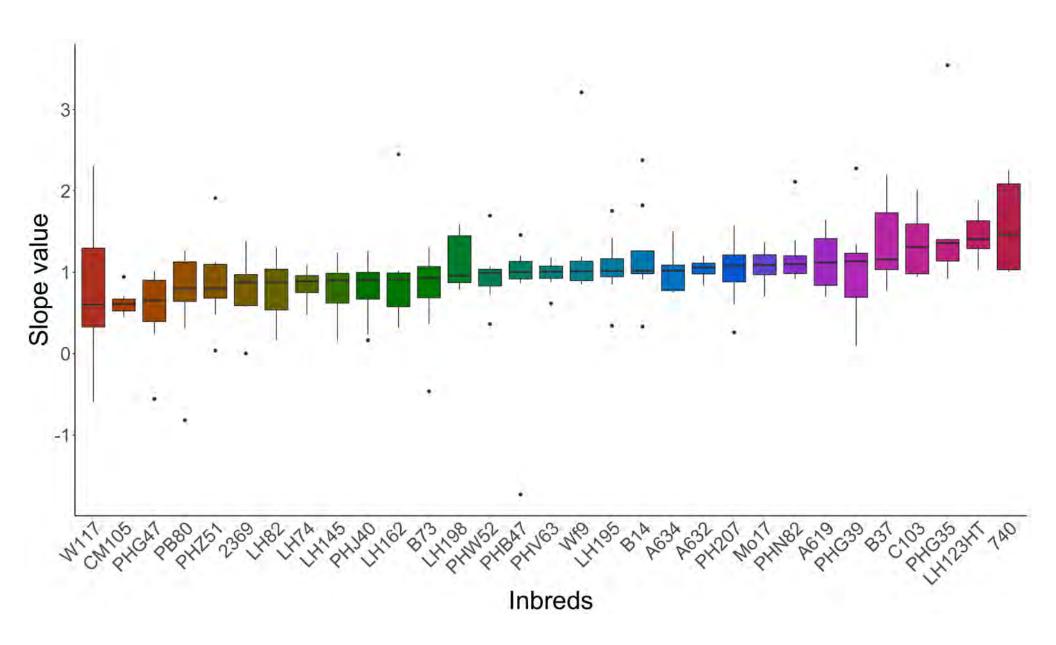


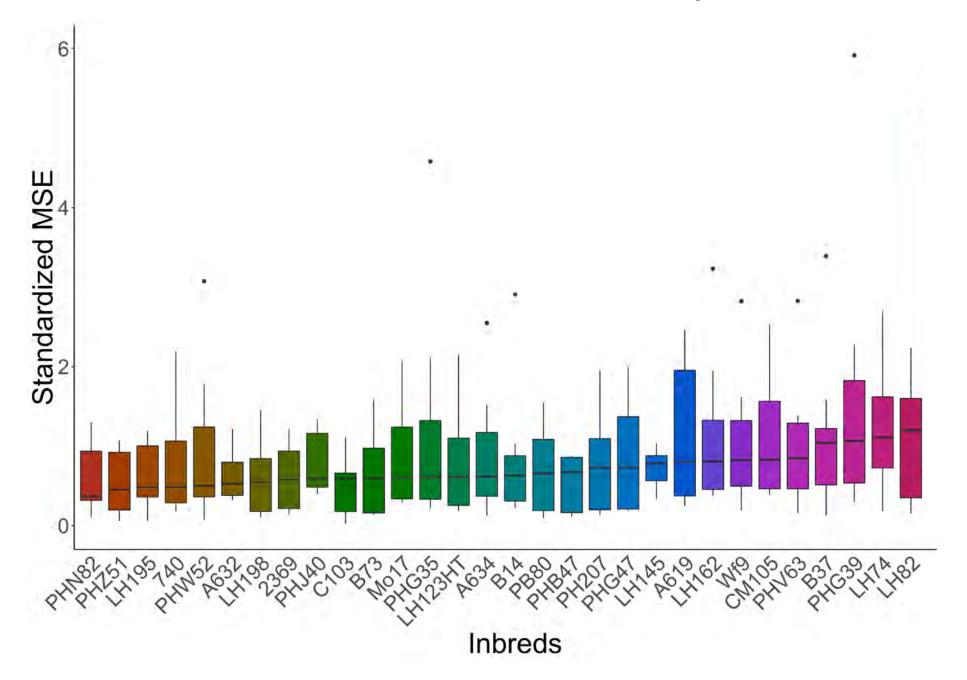
## What is the distribution of GxE among traits?



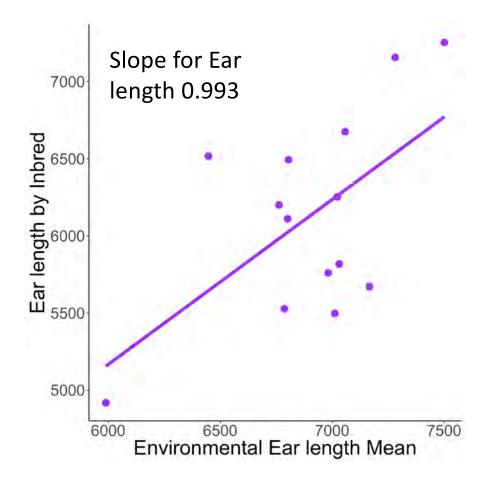
## What is the distribution of GxE among traits?



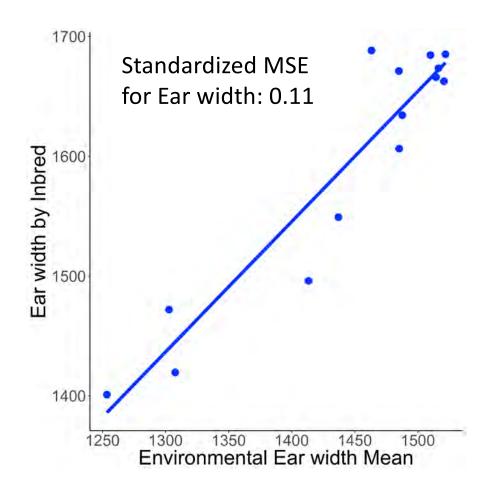


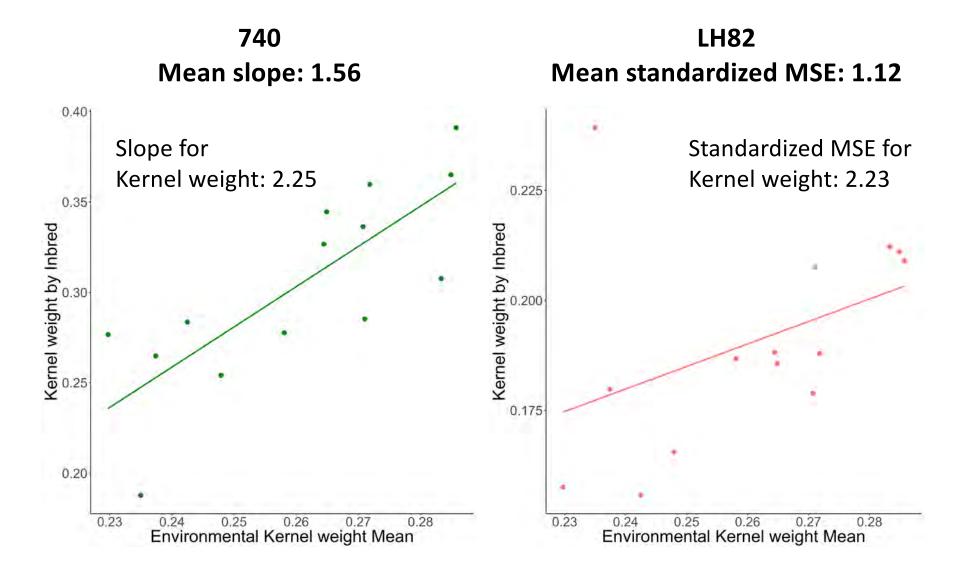


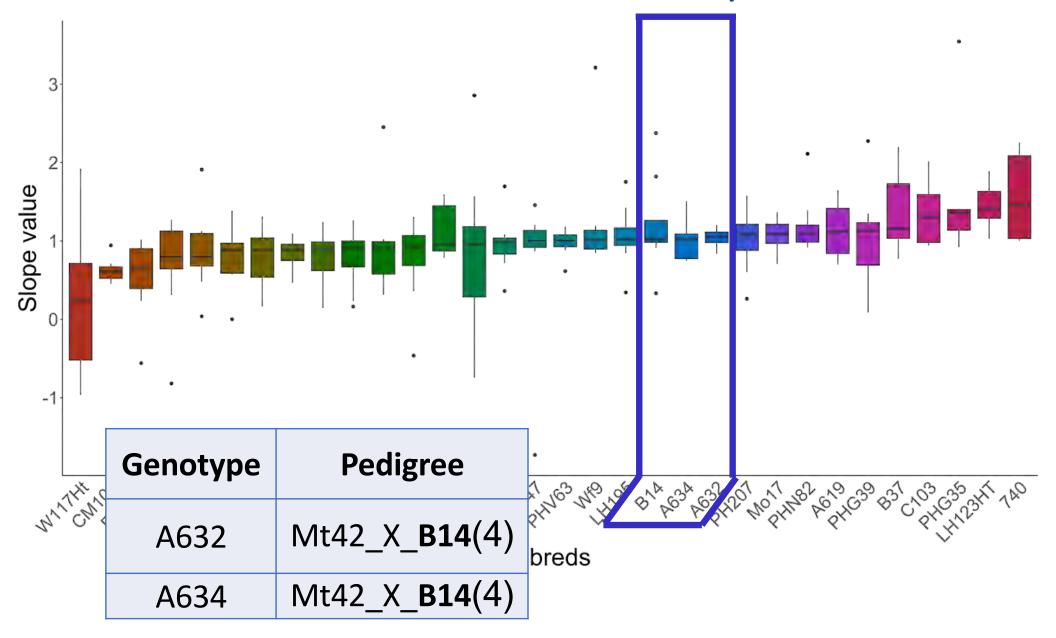
A634 Mean slope: 1.007

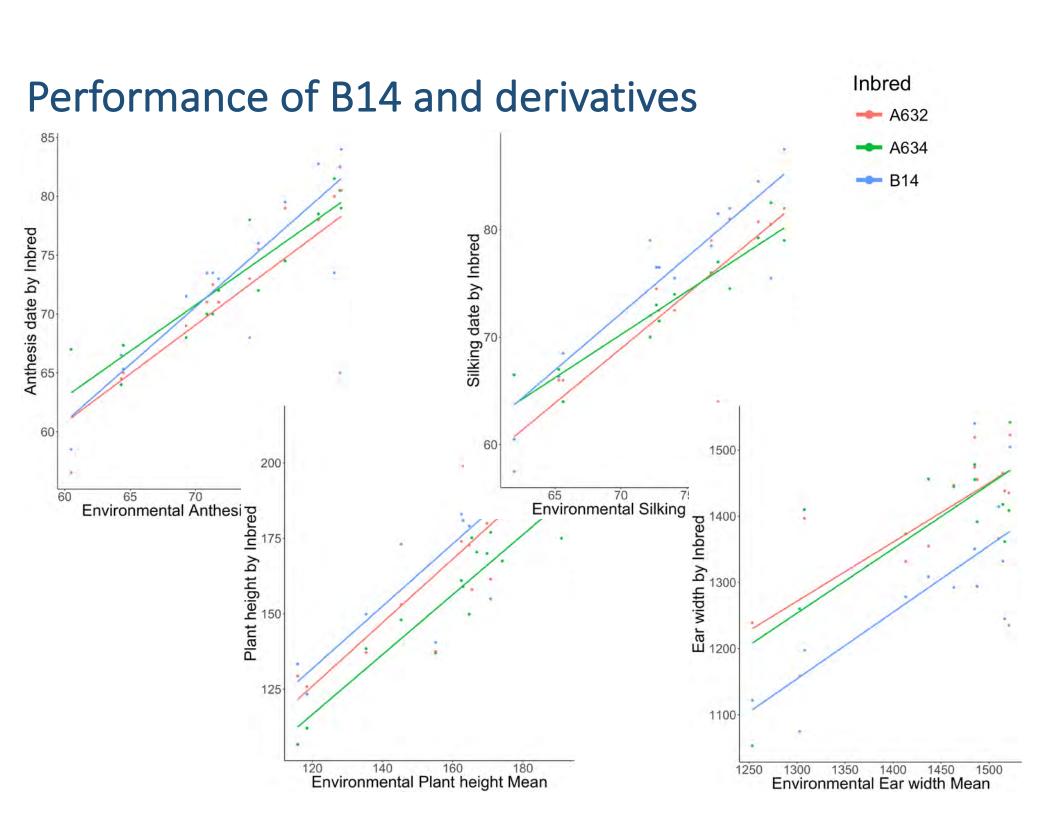


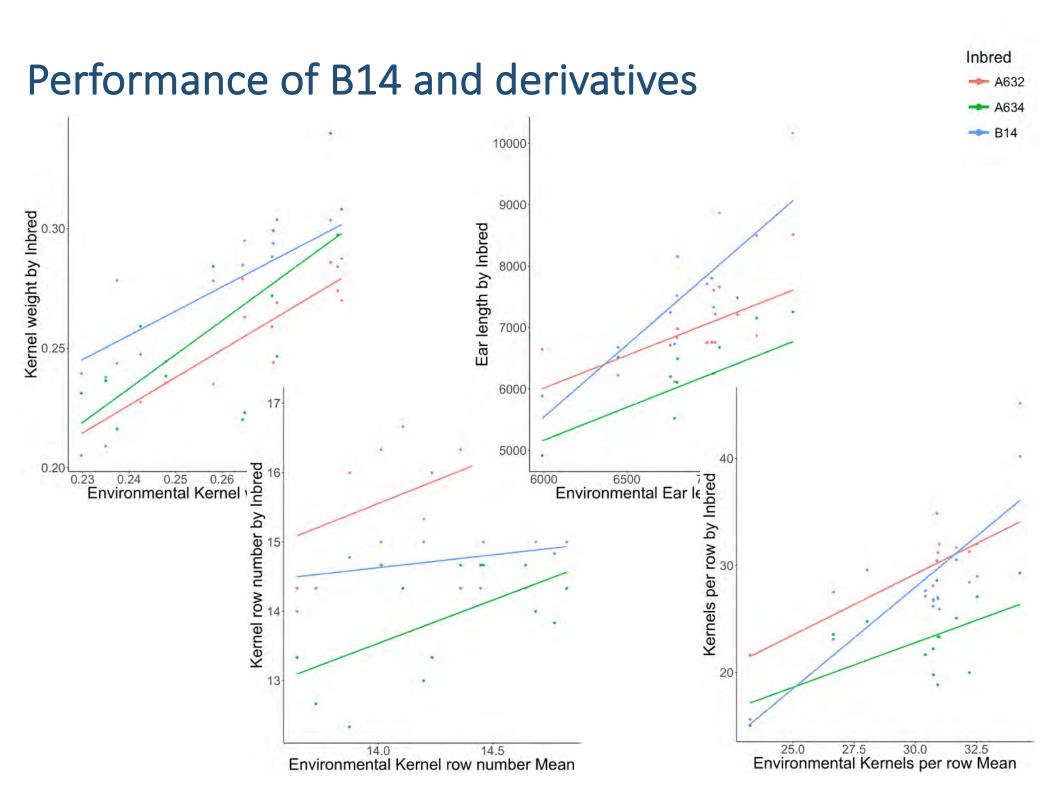
PHN82
Mean standardized MSE: 0.60



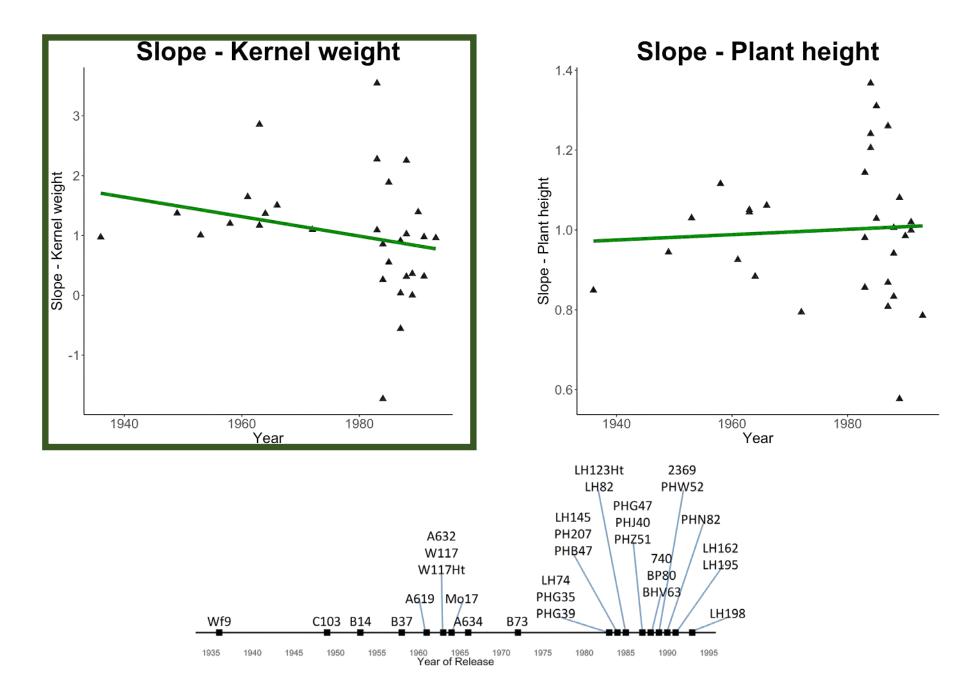




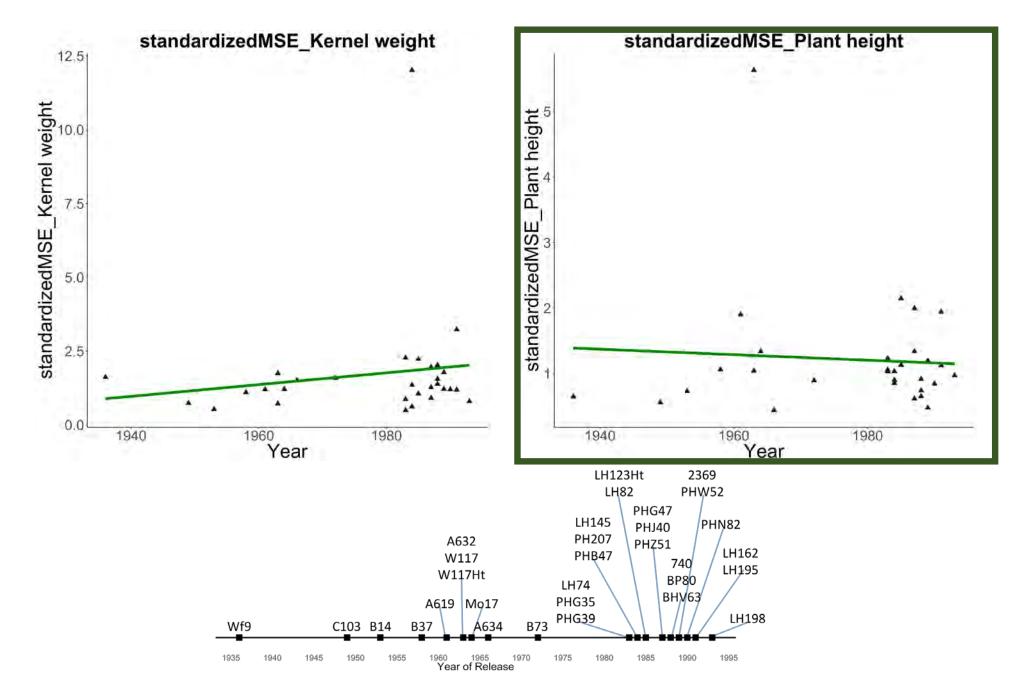




#### How does GxE change w.r.t. inbreds' year of release?



### How does GxE change w.r.t. inbreds' year of release?



#### Final remarks

- ♦G x E interaction contributes significantly to trait values, making it important to consider in predictions
- ♦ Ear/kernel traits showed greater GxE interaction—good "indicator traits" for future experiments
- ♦Some inbreds show better stability than others with exceptions for certain traits
- More recent inbreds show greater stability and improved predictability
- ♦ Much more to explore within this dataset and through incorporating the environmental data collected for each location

#### GxE Consortium: Data Usage Disclaimer

This presentation includes data analysis and interpretation conducted by the presenter and does not necessarily reflect the observations and conclusions of the GxE Consortium.

#### Acknowledgements

#### **♦**GxE 2014 Inbred Trial collaborators

- **♦** Jonathan Lynch
- **♦**Randy Wisser
- **♦Joe Knoll**
- **♦**Nick Lauter
- **♦** Pat Schnable
- **♦**Torbert Rocheford
- ♦ Sherry Flint-Garcia
- ♦Jim Holland
- **♦**Aaron Lorenz
- **♦Ed Buckler**
- ♦ Margaret Smith

- ♦ Rebecca Nelson
- **♦**Mike Gore
- **♦**Wenwei Xu
- ♦ Natalia de Leon
- ♦Shawn Kaeppler
- **♦**Candy Hirsch
- ♦ Nathan Springer
- **♦**Martin Bohn
- **♦**Seth Murray

## ♦ University of Wisconsin collaborators

- ♦ Nathan Miller
- **♦** Edgar Spalding
- ♦ Jane Petzoldt
- **♦Jonathan Renk**

#### G X E Cooperators

Principal Investigators who grew GxE trials in 2014-2016

- **Martin Bohn (UIUC)**
- **♦ Ed Buckler (ARS Cornell)**
- Ignacio Ciampitti (KSU)
- Jode Edwards (ARS ISU)
- **Sherry Flint-Garcia (ARS)**
- **Christopher Graham (SDSU)**
- Mike Gore (Cornell)
- **Candy Hirsch (UMN)**
- Jim Holland (ARS NCSU)
- **Elizabeth Hood (AR-State)**
- **David Hooker (Guelph)**
- Fiona Goggin (Univ AR)
- ♦ Shawn Kaeppler (UW)



AND ENVIRONMENTAL SCIENCES







**Judith Kolkman (Cornell)** 

- **Greg Kruger (UNL)**
- Nick Lauter (ARS ISU)
- Liz Lee (Guelph)
- Natalia de Leon (UW)
- Sanzhen Liu (KSU)
- **Argelia Lorence (AR-State)**
- Aaron Lorenz (UMN)
- Jonathan Lynch (PSU)

- Torbert Rocheford (Purdue)













**Oscar Rodriguez (UNL)** 

**Cinta Romay (Cornell)** 

James Schnable (UNL)

Rajandeep Sekhon (Clemson)

**Margaret Smith (Cornell)** 

Nathan Springer (UMN)

**Peter Thomison (OSU)** 

Mitch Tuinstra (Purdue)

Jason Wallace (UGA)

Randy Wisser (UDel)

Wenwei Xu (TAMU)

**Kurt Thelen (MSU)** 

Pat Schnable (ISU)

**Brian Scully (ARS)** 









- **Steve Moose (UIUC)**
- **Seth Murray (TAMU)**
- **Rebecca Nelson (Cornell)**













#### **Genomes To Fields Collaborators**

<sup>†</sup>G2F Executive Committee members \*GxE Coordinating Groups ~G2F co-lead

- Naser Alkhalifah (ISU)
- Martin Bohn (UIUC)\*
- Ed Buckler (ARS)†
- Darwin Campbell (ISU)\*
- Ignacio Ciampitti (KSU)
- James Clohessy (Cornell)
- Liang Dong (ISU)
- Jode Edwards (ARS)\*
- David Ertl (IA Corn)\*†
- Sherry Flint-Garcia (ARS)\*
- Joseph Gage (UW)
- Jack Gardiner (ISU)\*
- Fiona Goggin (Univ AR)
- Byron Good (Guelph)
- Mike Gore (Cornell)
- Christopher Graham (SDSU)
- Patricio Grassini (UNL)
- Jerry Hatfield (ARS)
- Candy Hirsch (UMN)
- Jim Holland (ARS)
- Elizabeth Hood (AR-State)
- David Hooker (Guelph)

- Diego Jarquin (UNL)\*
- Shawn Kaeppler (UW)†
- Joe Knoll (ARS)
- Judith Kolkman (Cornell)
- Greg Kruger (UNL)
- Nick Lauter (ARS)
- Carolyn Lawrence-Dill (ISU)\*
- Liz Lee (Guelph)\*
- Zhizhai Liu (TAMU)
- Natalia de Leon (UW)\*†~  $\diamondsuit$
- Alex Lipka (UIUC)
- Argelia Lorence (AR-State)
- Aaron Lorenz (UMN)\*  $\diamondsuit$
- Jonathan Lynch (PSU)†  $\diamondsuit$
- Nathan Miller (UW)
- Steve Moose (UIUC)
- Seth Murray (TAMU)\*
- Rebecca Nelson (Cornell)
- Torbert Rocheford (Purdue)
- $\diamondsuit$ Oscar Rodriguez (UNL)
- Cinta Romay (Cornell)\*  $\diamondsuit$
- Emily Rothfusz (UW)\*

- James Schnable (UNL)
- Pat Schnable (ISU)+~
- Brian Scully (ARS)
- Rajandeep Sekhon (Clemson)
- Kevin Silverstein (UMN)
- Margaret Smith (Cornell)
- Edgar Spalding (UW)
- Nathan Springer (UMN)†
- Srikant Srinivasan (ISU)\*
- Yiwei Sun (ISU)\*
- Kurt Thelen (MSU)
- Peter Thomison (OSU)
- Kelly Thorp (ARS)
- Mitch Tuinstra (Purdue)
- Jason Wallace (UGA)
- Renee Walton (ISU)
- Rod Williamson (IA Corn)
- Randy Wisser (UDel)\*
- Wenwei Xu (TAMU)
- Cheng-Ting Yeh (ISU)
- Jianming Yu (ISU)



AND ENVIRONMENTAL SCIENCES









JNIVERSITY





























#### **Genomes To Fields Sponsors**























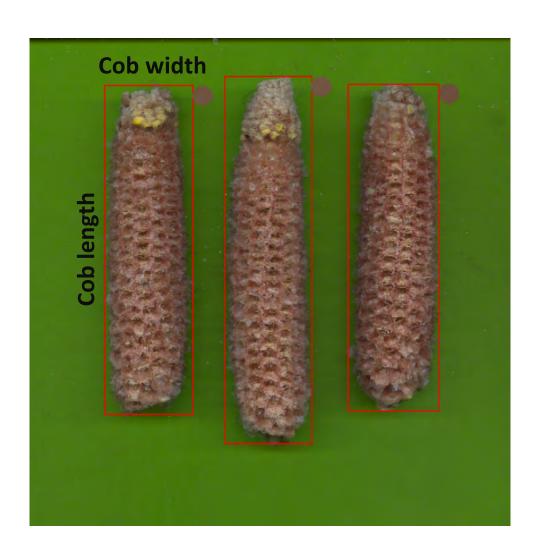




Thank you for your attention!

Any questions?

## **Imaging Output: Cobs**



#### References

- ♦ Bernardo, R. 2010. Breeding for Quantitative Traits in Plants. Second ed. Stemma Press, Woodbury, Minnesota.
- → Miller, N.D., N.J. Haase, J. Lee, S.M. Kaeppler, N. de Leon, and E.P. Spalding. 2016. A robust, high-throughput method for computing maize ear, cob, and kernel attributes automatically from images. Plant J. Available at http://doi.wiley.com/10.1111/tpj.13320.