



UNIVERSITY OF WISCONSIN-MADISON

How Do Advances in Corn Breeding Improve the Corn for Silage?

World Dairy Expo 2017

Dairy Forage Seminar Stage

Natalia de Leon – Department of Agronomy – University of Wisconsin

October 5th, 2017



Outline

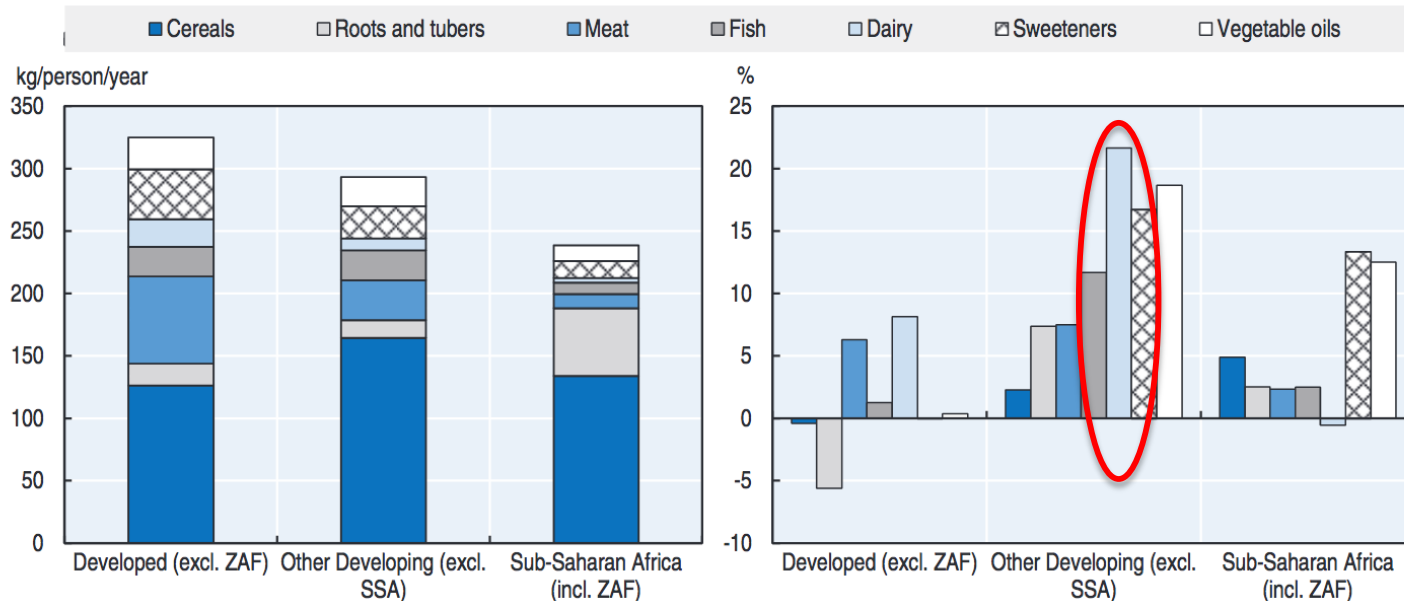
- ✧ Silage Breeding and the UW Program?
- ✧ Tools to Increase Breeding Outcomes
- ✧ Future Challenges

Demand for Silage?

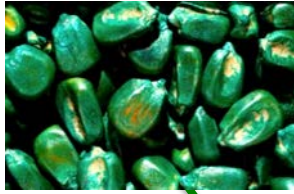
- ✧ While income growth in emerging economies is projected to be weaker, increased global population is expected to increase demand for dairy products

Figure 1.2. **Per capita food consumption by region**

Kg/cap/year in 2025 (left) and growth 2025 vs. 2013-15 (right)



Seed to Seed: 800X Biomass Return



~30 kg seed/ha

~75 days



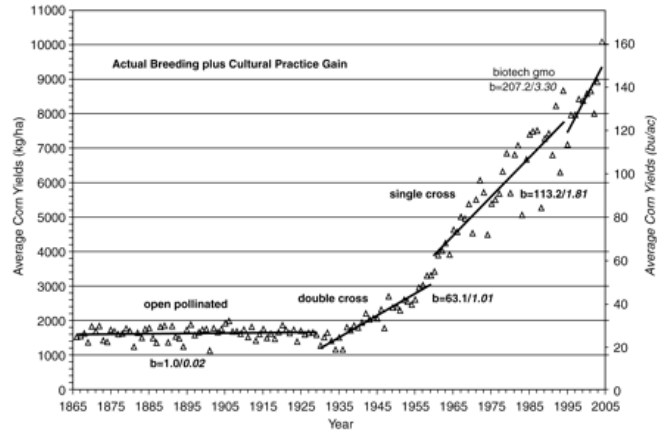
~12 Mg/ha
vegetative biomass
at flowering

~45 days



~12 Mg/ha
grain

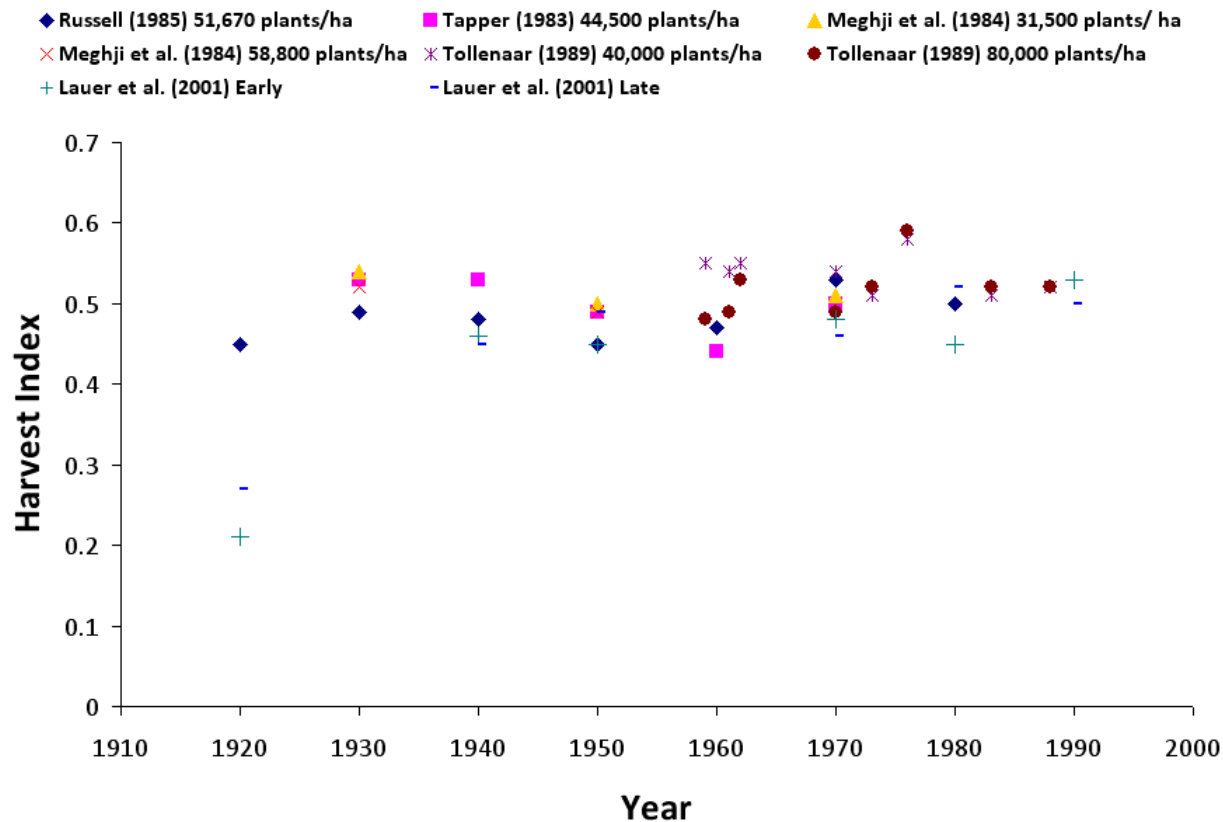
Relative Contribution of Grain vs Stover



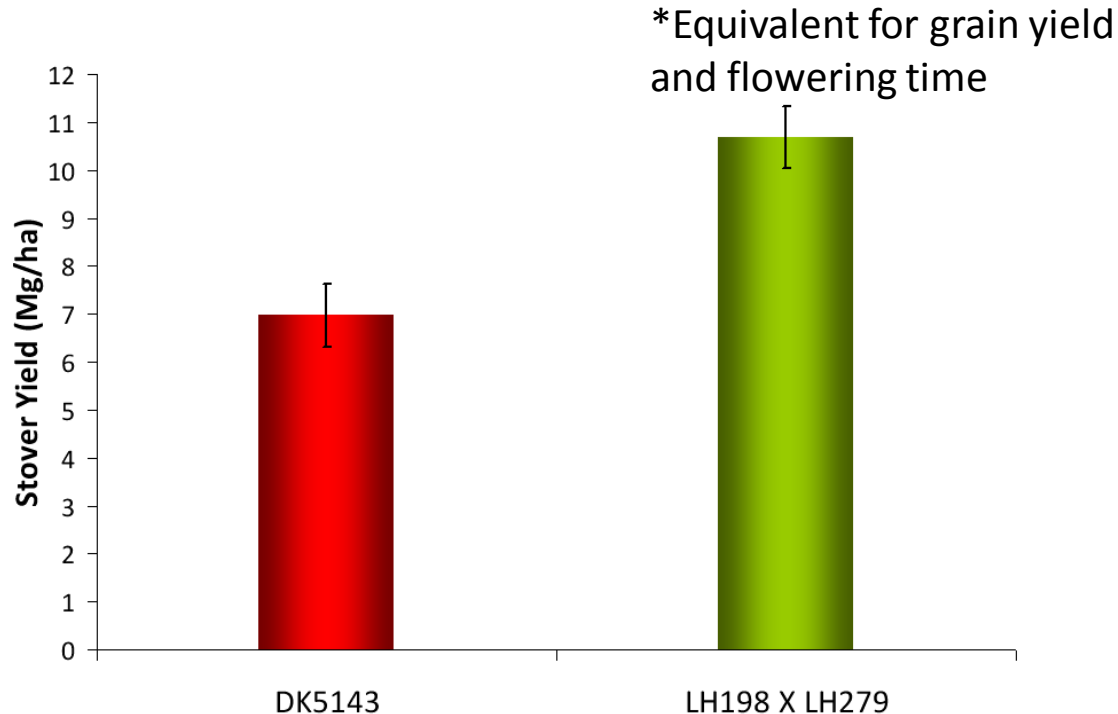
VS



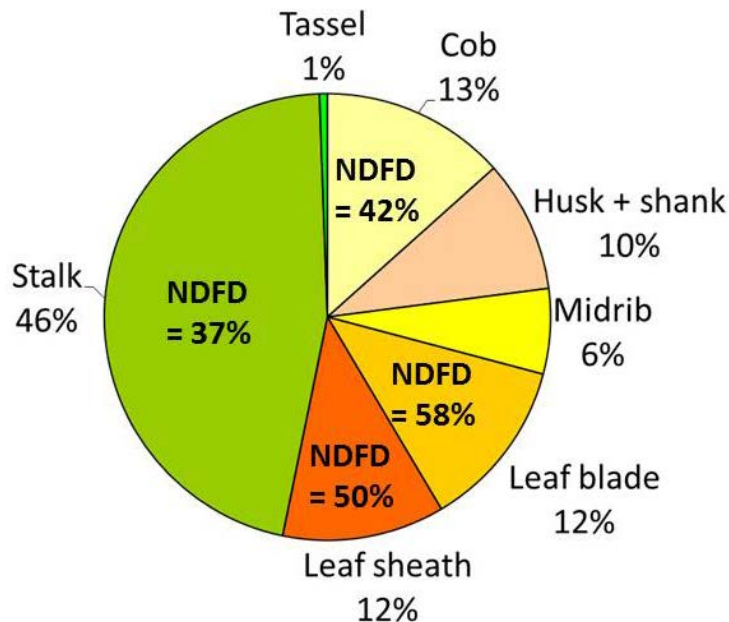
Has Harvest Index Remained Constant in Corn?



Variation for Stover Yield Among Elite Grain Hybrids



Where is Biomass Quality Coming From?



❖ Comparison:

– *bm3* (=COMT) vs
isoline ~10 to 20%
increase in NDFD

– Increase blade sheath
by 50% ~5% increase in
NDFD

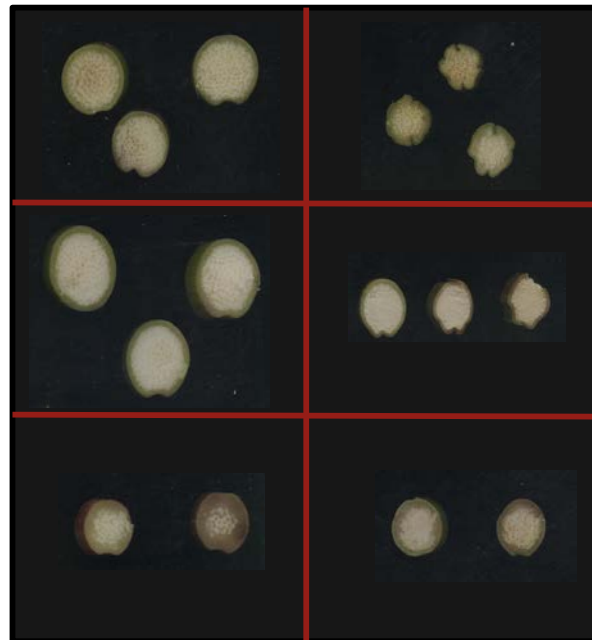
Relationship between Digestibility and Anatomical Traits



Genotypes with
low sugar release



Genotypes with
high sugar release



Silage Production and the UW Program

- ✧ Approximately 7% of the corn acreages harvested in the U.S. in 2016 were silage acres and WI is the largest producer with ~790,000 acres
- ✧ The UW program focuses on the development of corn varieties with enhanced compositional attributes and forage yield
 - ❖ High yield
 - ❖ High energy (high digestibility)
 - ❖ High intake potential (low fiber)
 - ❖ High protein
 - ❖ Proper moisture at harvest for storage





Scope of the Program


- ✧ Between 2,000 and 3,000 evaluation plots are dedicated to the silage breeding program yearly
- ✧ Evaluation includes two main locations – West Madison and Arlington, WI
- ✧ Advanced lines are evaluated by the UW Extension Program at an additional 3 to 4 location based on relative maturity
- ✧ Additional silage plots are dedicated to silage research activities which varies from year to year
- ✧ Release more than 8 breeding populations, UW NIRS silage quality prediction equations and 25 inbred lines since inception

Selection Criteria – MILK 2006

Milk/ton	Milk/acre
lbs	lbs

2005 EXPT WQS198 - Forage yield and nutritional evaluation (Madison and Arlington)

05No.	Entry	Dry matter	Yield	NDF	IVD	IVNDFD	Protein	Starch	Milk/ton	Milk/acre
		%	t/a	%	%	%	%	%	lbs	lbs
1	12004-1-4 X LH198	44.3	7.56	46.3	80.6	58.1	7.94	29.9	3244	24560
4	12021-3-4 X LH198	41.0	6.33	46.3	81.2	59.4	7.90	28.2	3258	20642
6	12028-3-1-5-1 X LH198	44.0	7.77	46.9	80.4	58.0	7.89	29.2	3225	25049
8	12028-3-2 X LH198	43.3	8.27	47.8	80.6	59.3	8.10	29.0	3224	26641
9	12028-3-3 X LH198	45.0	8.41	45.7	80.4	57.1	7.91	31.7	3242	27231
10	12028-3-4 X LH198	46.3	7.61	49.0	78.8	56.7	7.70	28.3	3118	23743
13	12057-1-1-2 X LH198	42.3	7.47	47.4	80.0	57.8	7.95	29.1	3199	23912
15	12057-1-5 X LH198	43.8	8.99	43.9	80.8	56.1	7.74	33.5	3281	29524
20	12135-3-1-3-1 X LH198	46.5	8.23	47.1	79.8	57.1	7.59	30.7	3192	26314

Milk 2006


University of Wisconsin Corn Silage Evaluation System
Randy Shaver, Dept. of Dairy Science *Patrick Hoffman, Dept. of Dairy Science*
Joe Lauer, Dept. of Agronomy *Jim Coors, Dept. of Agronomy*
Sample values entered here must correspond to lab average and incubation times information entered in "UserInputGuide" Worksheet cells G25 and G27.

Critical Data Entry
Required Inputs
Calculated Outputs

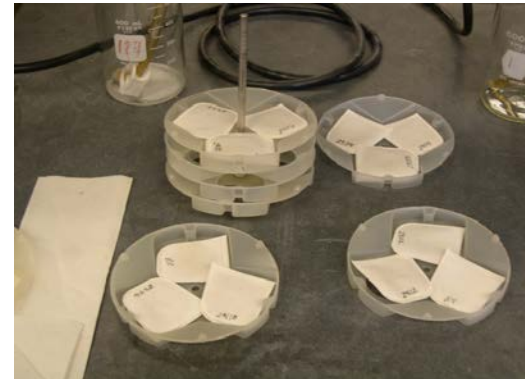
Field ID	Lab ID	Processed	Kernel	KPS	DSA	IS-IV	DM	CP	NDF	IVD	Starch	Ash	Fat	DM Yield	TDN-1x	IE-3x	Calculated Milk per Ton Index	Calculated Milk per Acre Index
		%	%	%	%	%	% DM	% DM	% DM	% DM	% DM	% DM	% DM	tons/acre	% DM	Meals/ton DM	lb/ton DM	lb/acre
"normal"	L001	no					35.0	8.8	45.0	59	27.0	4.3	3.2	7.0	69.7	6.68	3169	22182
1	L002	yes					35.0	8.8	45.0	59	27.0	4.3	3.2	7.0	71.0	6.70	3273	22910
							-	-	-	-	-	-	-	-	#VALUE!	#VALUE!	#VALUE!	#VALUE!
							-	-	-	-	-	-	-	-	#VALUE!	#VALUE!	#VALUE!	#VALUE!

Quality Analysis Methods

✧ **Neutral detergent fiber (NDF); Acid detergent fiber (ADF); Acid detergent lignin (ADL) and In vitro true digestibility (IVTD)** following Goering and Van Soest, 1970 with modifications (ANKOM system)

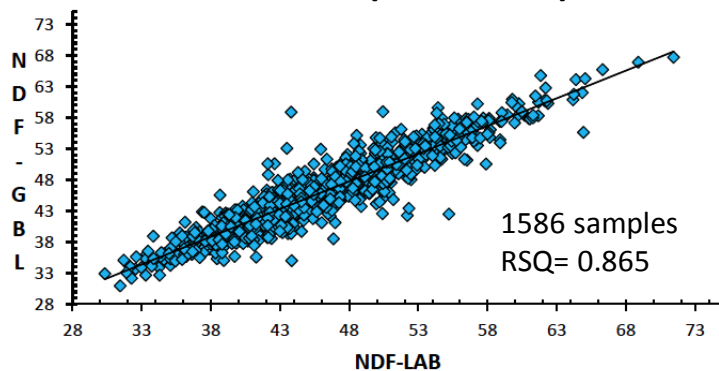
✧ **Protein** = Measure nitrogen using the Leco FP-528 nitrogen analyzer (N X 6.25)

✧ **Starch** = Predicted using whole plant silage global NIRS equation

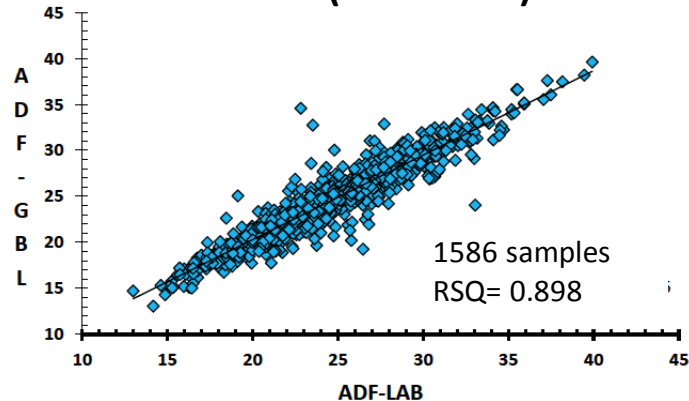


2016 NIRS Global Equation Calibration

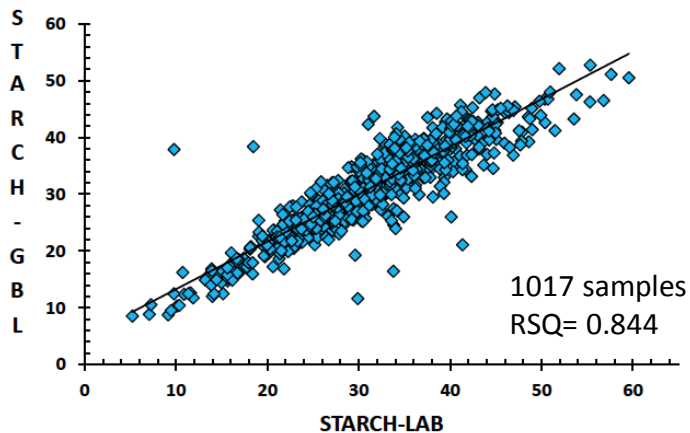
2016 NDF (GBL – LAB)



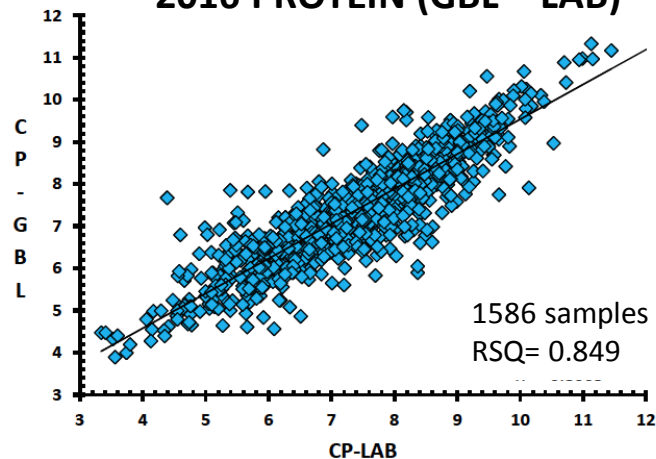
2016 ADF (GBL – LAB)



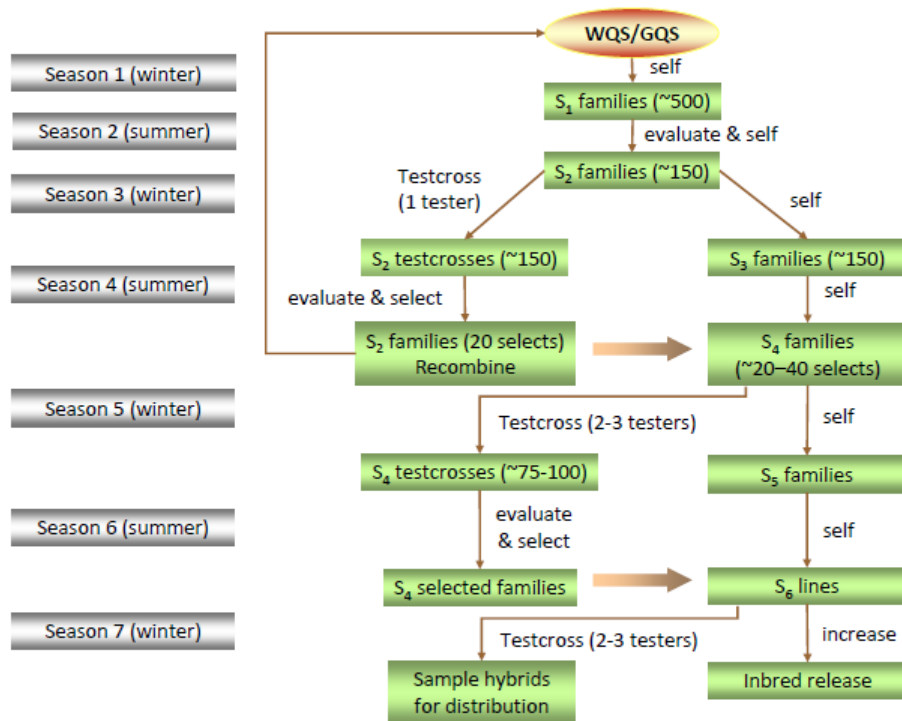
2016 STARCH (GBL – LAB)



2016 PROTEIN (GBL – LAB)



Selection Protocol

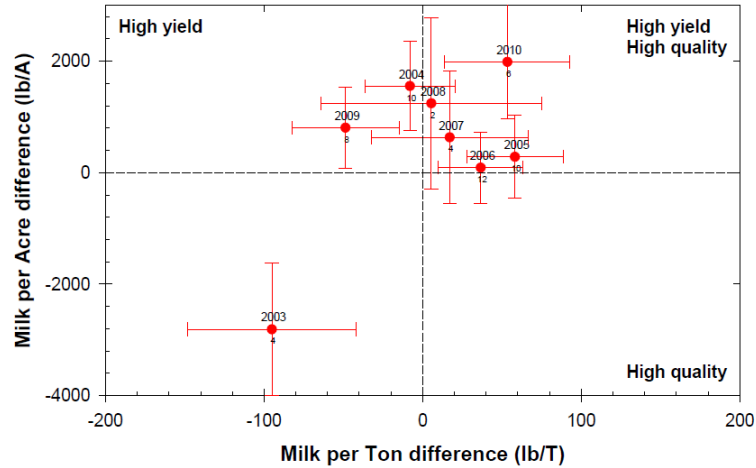


WQS - Broad-based high quality synthetic (low fiber, lignin and silica) then crossed by H99 and Mo17 (100 to 110RM)

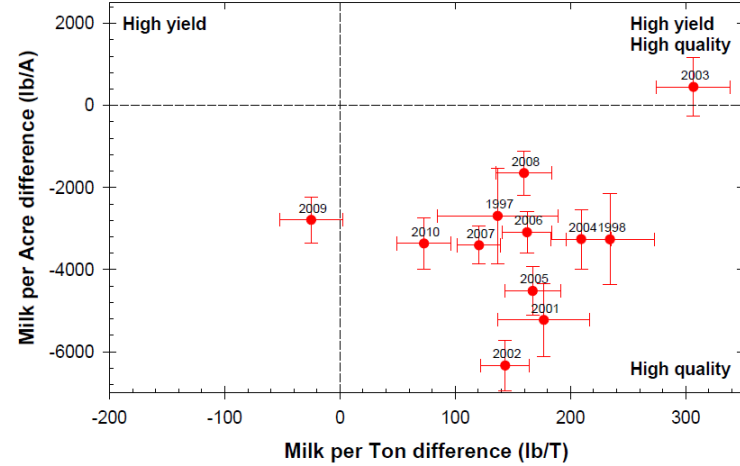
GQS - CUBA164:S1517 and CUBA117:S1520 populations are from the Stiff Stalk Synthetic background (75%)

Also germplasm from the Germplasm Enhancement of Maize (GEM) program

WQS Germplasm



WQS derived lines



***brown midrib3* lines**

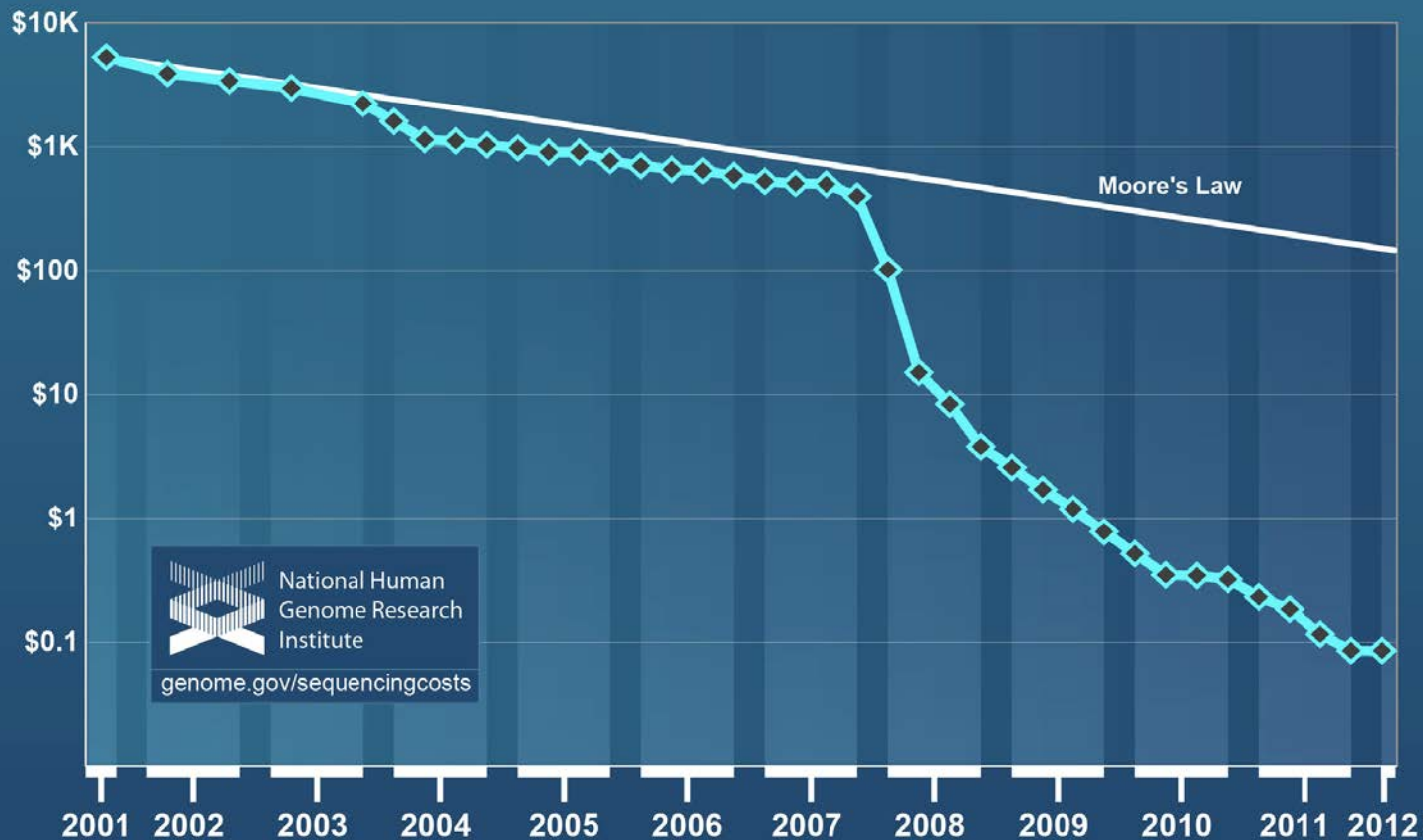
Joe Lauer - Data from UW Extension - Corn Trial

Outline

- ✧ Silage Breeding and the UW Program?
- ✧ Tools to Increase Breeding Outcomes
- ✧ Future Challenges



Cost per Raw Megabase of DNA Sequence



Exploiting Genetic Variability

Average height = 100

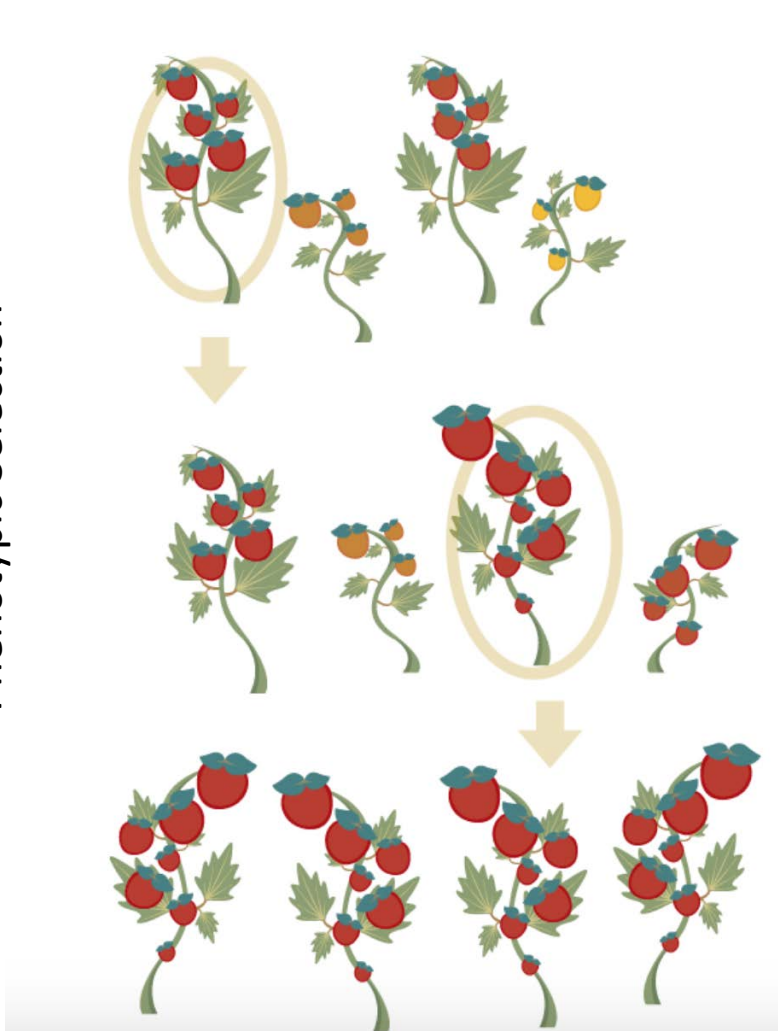
C T G **T** T A T

Average height = 50

C T G **A** T A T

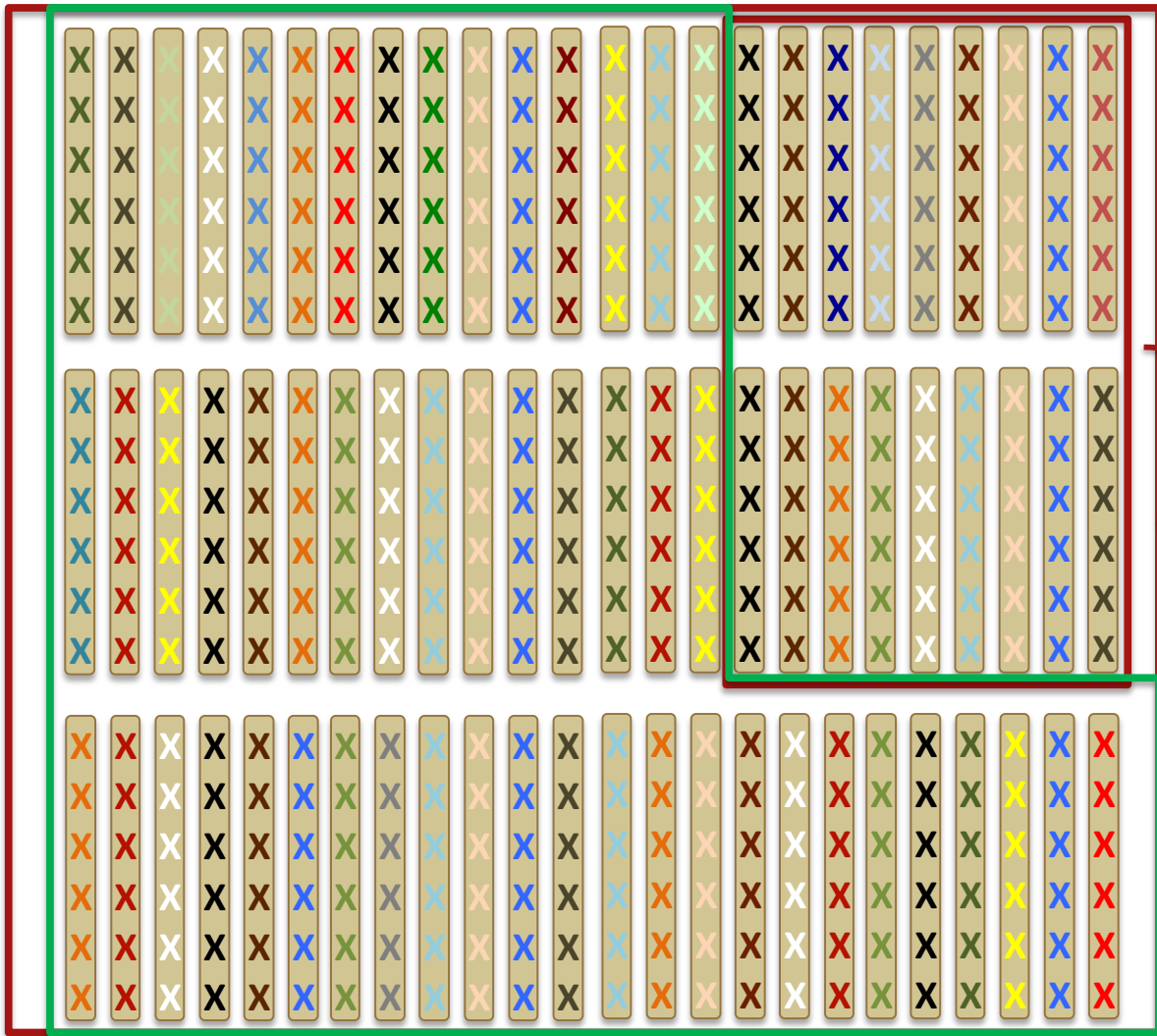


Phenotypic Selection



Genomic Selection





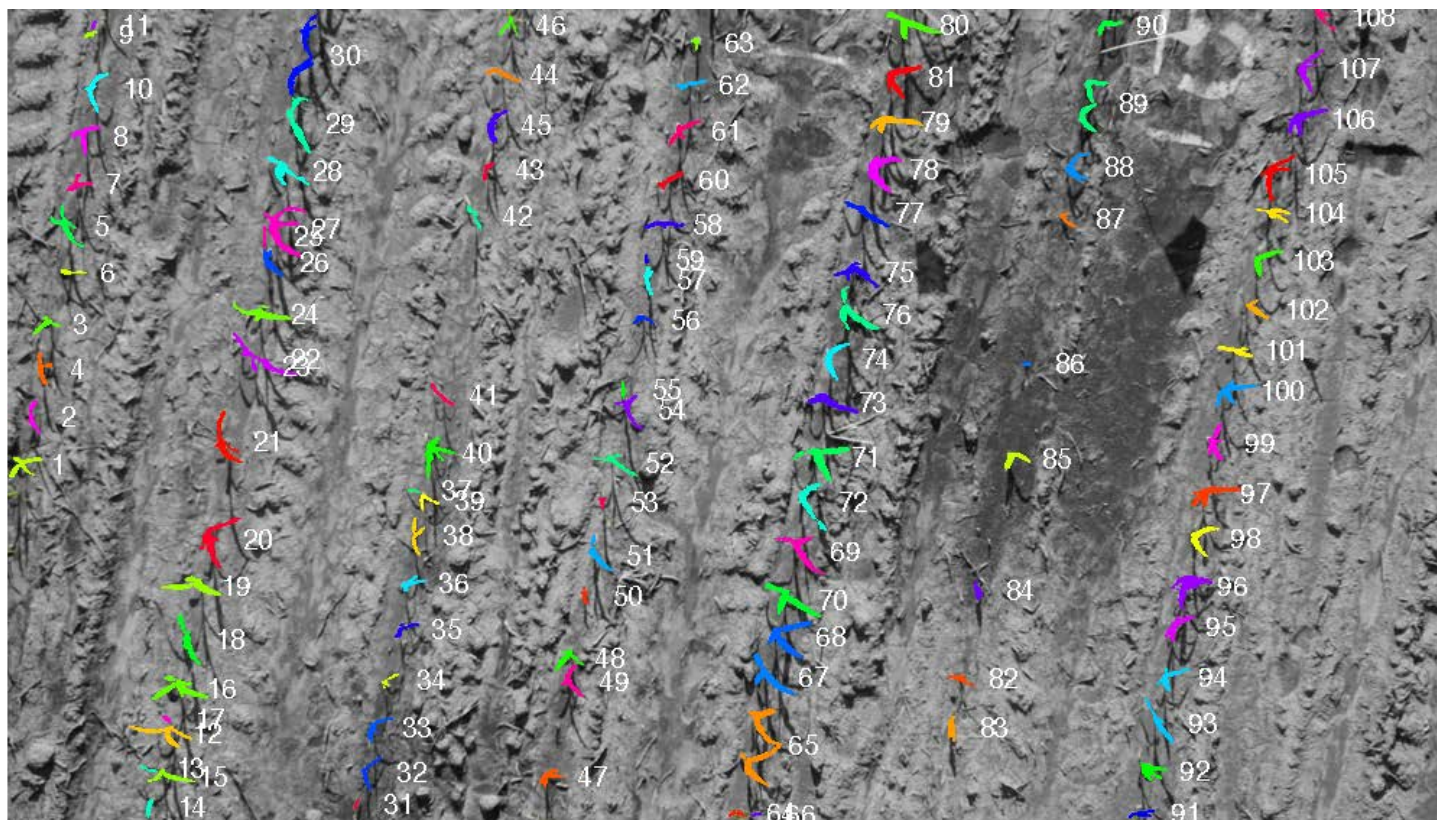
$$y_i = \mu + \sum_k \beta_k x_{ik} + e_i$$

Silage includes many traits

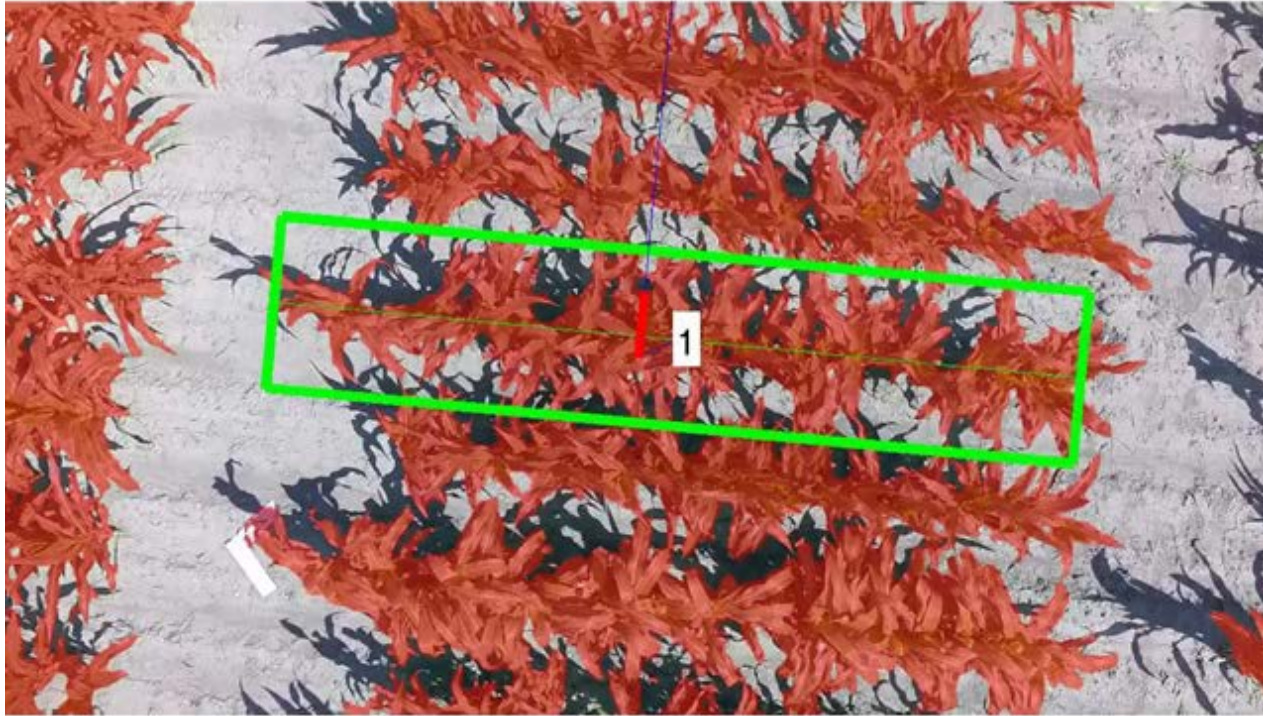
Phenotyping



Stand Counts



Growth Rate and Tasseling



Nathan Miller
Spalding Lab

Outline

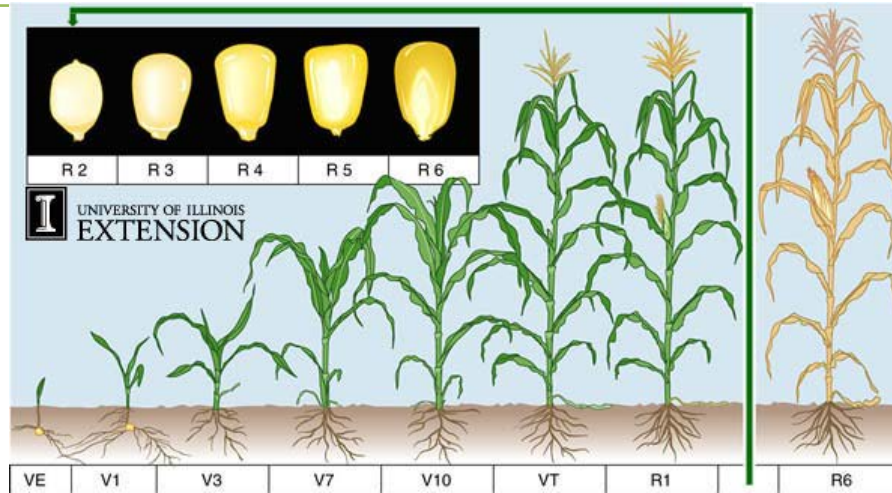
- ✧ Silage Breeding and the UW Program?
- ✧ Tools to Increase Breeding Outcomes
- ✧ Future Challenges

Phenotypes are Modified by the Environment over Time

Factor	VE	V6	V12	V18	R1	R6
Frost (< 28 F)	0	100	100	100	100	0
Hail (max)	0	53	81	100	100	0*
Drought/Heat (%/day)	--	--	3	4	7	0
Flooding (<48 h)	severe	severe	0	0	0	0

* Assuming no ear dropage

<http://corn.agronomy.wisc.edu/Management/L011.aspx>





Day 1

Day 2

Day 3

Day 4

Day 5



Edgar Spalding
UW Madison

What Our Eyes Cannot See



Take Home Message:

- ✧ Need for varieties with improved silage characteristics is expected to continue to increase due to international demand
- ✧ Equipment and management tools have helped research
- ✧ Genotypic information is cheap, meaningful, high-throughput phenotyping is the bottleneck
- ✧ Interdisciplinary developments are needed to increase efficiency of phenotyping (current bottleneck)
- ✧ The UW Corn Silage Breeding Program continues to serve as a source of germplasm, research and a tool for training of students

Acknowledgements:





THANK YOU.

- ✧ Shawn Kaeppler and UW Group
- ✧ Edgar Spalding group – UW Madison
- ✧ Robin Buell group – Michigan State
- ✧ Candy Hirsch – Univ of MN

