

Breeding programs to mitigate climate change and environment pressures on crops

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UPOV - Seminar on the role of plant breeding and plant variety protection in enabling agriculture to mitigate and adapt to climate change

October 11, 12, 26; 2022

File under seed com & D. Saton A. Requests ames, Down march 7 , 1926

HA. Wallace Des Moines Jana

Dear Mr. quallans at the corn show last February, you said that if I would write to you along is mand you might be able to let me have some inbred corn to cross and enter in the Com yield contest. So & an writing you in hopes that you will have some to spare. I are a sophomore Farme, Crafe, and Soils stadent at ames, so naturally very much interested in com. De it best when crossing com to use the paper sach method, do no to creater on both parents; or just cart ile Tascelo aff of one strain? I am going to school at ames, but would want any corr sent home to Blaconsfield, Jown. you an eitles and it C.O.D or lit me know, what the postage is. Hanking you in advance, I remain, Jamo Trakes

contrast and and

March 10, 1926.

Mr. H.A. Wallace DesMoines, Iowa

Dear Mr. Wallace:

Our corn is doing fine, and I am having a lot of fun pulling out the tassles. I have already pulled out about half the tassels. Some of the mother parents are so tall(over 8 ft) that I need a step-ladder to reach the tassels.

Beaconsfield, Iowa July 15, 1926

I think I can send you sees inbred strains of corm all right, but before sending them I would like to know just what kind of a plot you have. Do you have a half acre which is at least 500 feet away from any other corm, where there would not be much chance for pollen from other corm coming in on it? If so, I would send you one sort to use as a male parent and thirty or forty other sorts to use as fecale parents. I would want you to plant about twenty hills each of these differents female parents. We would then take four pounds of anch of these different combinations and enter them in the Jown Oorn Yield Contest in the spring of 1927. We can enter these coubingtions in the name of Baker and Wallace if you so desire.

If you take on this project, you will of course have to arguings to plant the corn with some care so you know in which row each of the different strains is planted, and then in July you will have to arguings to pull the tassais out every day, and in late September or early Ootsber you will have to harvest the different norts and label them. It is guite a little job.

Frite me further on the matter as to just her you want to go absad with this proposition. If you happen to be coming down to bes Molnes at any time , let me know in advance and drop around to the office.

HAWallace NN

Mr. Saymond Baker.

G.D. Station A. Ames, lows.

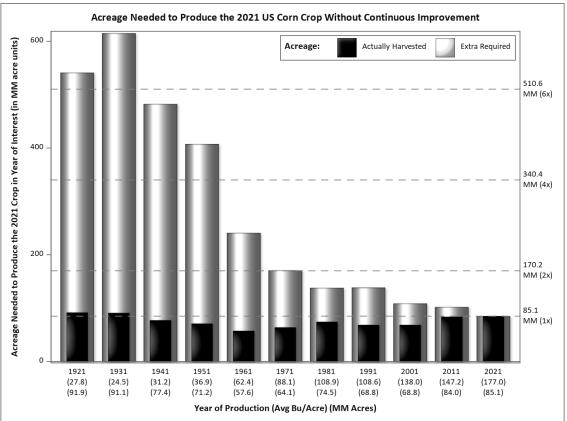
Dear Mr. Bakert

Sincerely yours,



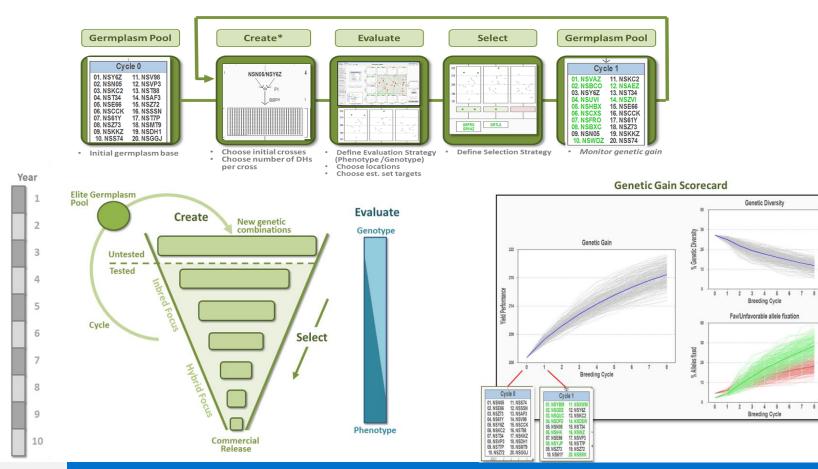
RAYMOND BAKER

Acreage impact of corn breeding and improved management practices





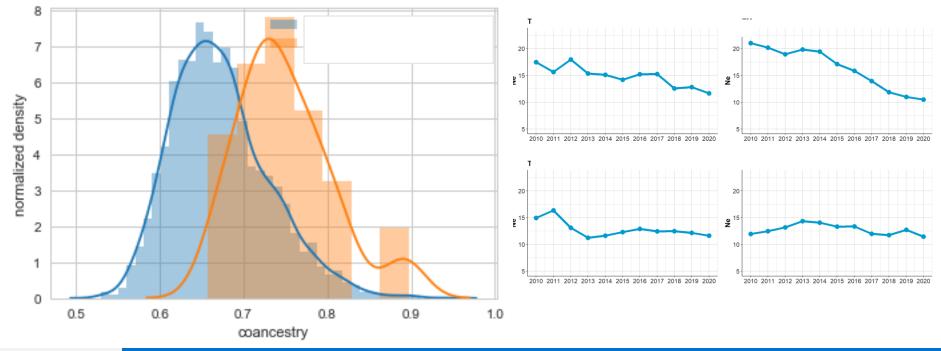
The Breeding Pipeline - Conduct pre-commercial product testing in target environments for multiple years





Genetic variation – trends over time

- Genetic variation is essential to achieve breeding goals
- Co-ancestry based pedigree and/or genotype
- Allelic diversity driven by population sizes in breeding programs





Maize Breeding – highly complex trait selection needs Genetic gain for any of these traits requires favorable variation and response to long-term selection, adding patience and adjusting breeding goals as the environmental pressures change over time

Agronomics

Yield

Test weight Grain moisture Grain dry-down Stalk lodging **Root lodging Plant height** Ear height **Brittle snap Drought tolerance** Emergence Stand establishment Early growth Cold tolerance

Pest Resistance

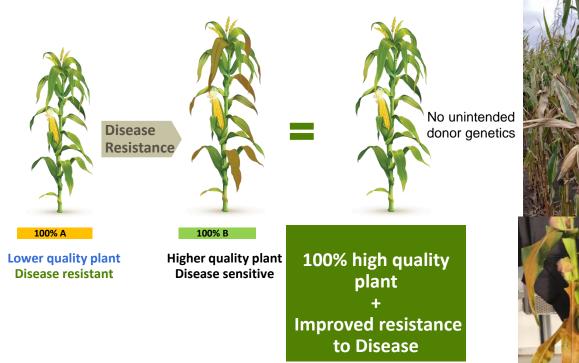
Gray leaf spot Northern leaf blight Southern leaf blight Stewart's wilt Rusts Smuts Anthracnose Diplodia Giberella Fusarium Diplodia Bacterial wilt

End-Use Traits

Mycotoxin Production in Grain Starch, Protein, & Oil Extractable starch Total fermentables Gross energy Digestible energy Food-grade Traits Silage quality traits



Native Genetics and CRISPR approach to Disease Control



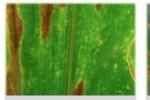




Accelerating Native Genetics for Disease Control

Plant Disease is a Major Challenge for Growers

In 2021, North American corn growers lost more than 318 MM bu¹ due to:





Northern Leaf Blight

Southern Rust

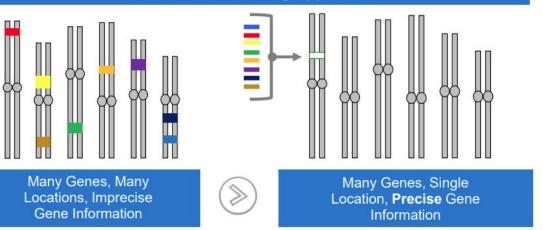


Gray Leaf Spot



Anthracnose Stalk Rot

Our Patent-Pending Approach



- Builds off germplasm advantages and improves genetic gain
- New breeding techniques unlock additional power of native genetics
- Multiple disease targets

- Multiple native genes for each disease target improves resistance and adds durability
- Simplified genetics assembled through gene editing accelerates plant breeding



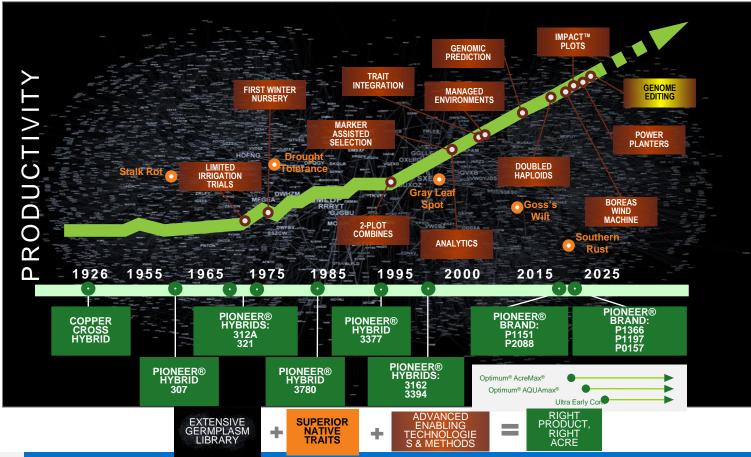


Genome editing and future potential – what if we could...go beyond plant breeding techniques and exceed current range of variation

- Chromosomal rearrangements (CR's) and potential to change phenotypes and "unlock" genetic variation (<u>https://www.nature.com/articles/s41477-020-00817-6)</u>
- Effective control of genomic recombination elements
- Enable co-location of native traits, unleashing major portions of the genome for improved maintenance of genetic diversity and additional favorable trait selection
- Multiple and simultaneous edits across numerous traits



Will Genome Editing reside on this chart in 2070?





Key Points

- Conduct plant breeding and pre-commercial product testing in the target environments for multiple years
- Genetic variation is essential to achieve breeding goals and mitigate climate change
 - Account for inevitable diversity decline over time
 - Leverage science and technology to create new favorable variation
- Plant breeding requires long-term selection, patience and adjusting breeding goals as the environmental pressures change over time
- Genome editing methods have potential for creating additional and needed variation to accomplish future environmental needs to feed a growing population
- Increased progress to minimize or eliminate biotic and abiotic stresses enables increased heritability and selection efficiency for grain yield improvement



Thank you!



