



...seeding a sustainable future



New transcription factor based technology & tools for future crop improvement

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Mendel Biotechnology, Inc.**

**New Phytologist Tansley Symposium
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Timberline Lodge, Mt. Hood, Oregon USA**

The challenge



John Stringer & Linda Horton

**we need to improve
crop productivity!**

Ag. Biotechnology Product Timelines

First generation (mid 1990s)

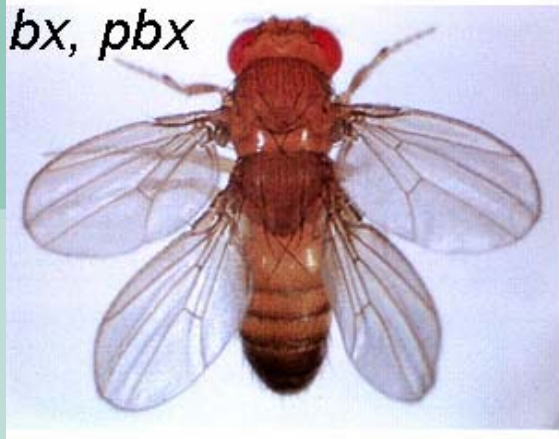
- single gene traits
- herbicide and insect resistance

Next generation (early 2010s)

- complex polygenic traits
 - e.g. yield and stress
- Based on whole pathway regulation

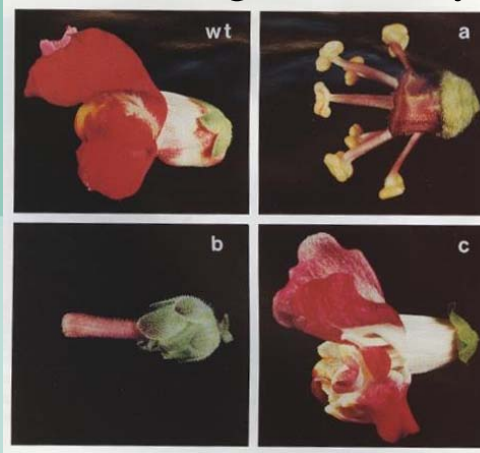
Body plan patterning

bx, pbx



e.g. Homeodomain proteins
McGinnis et al., 1984, Nature 308

Floral organ identity



e.g. MADS domain proteins
Coen & Meyerowitz, 1991, Nature 353

Floral symmetry



e.g. CYCLOIDEA-related proteins
Luo et al., 1996, Nature 383

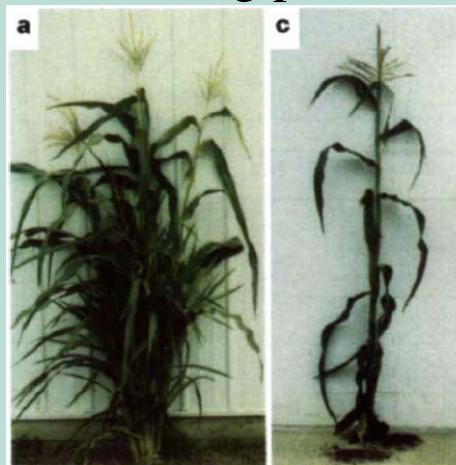
Transcription factors: master regulators of genetic pathways

Flowering time



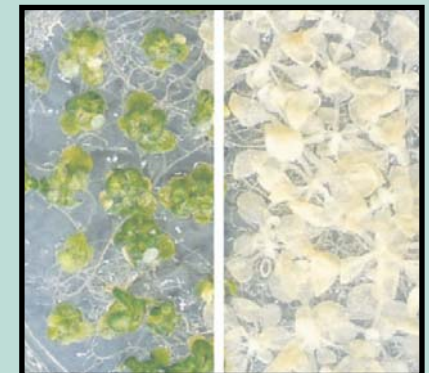
e.g. CONSTANS
Putterill et al., 1995, Cell 80

Branching pattern



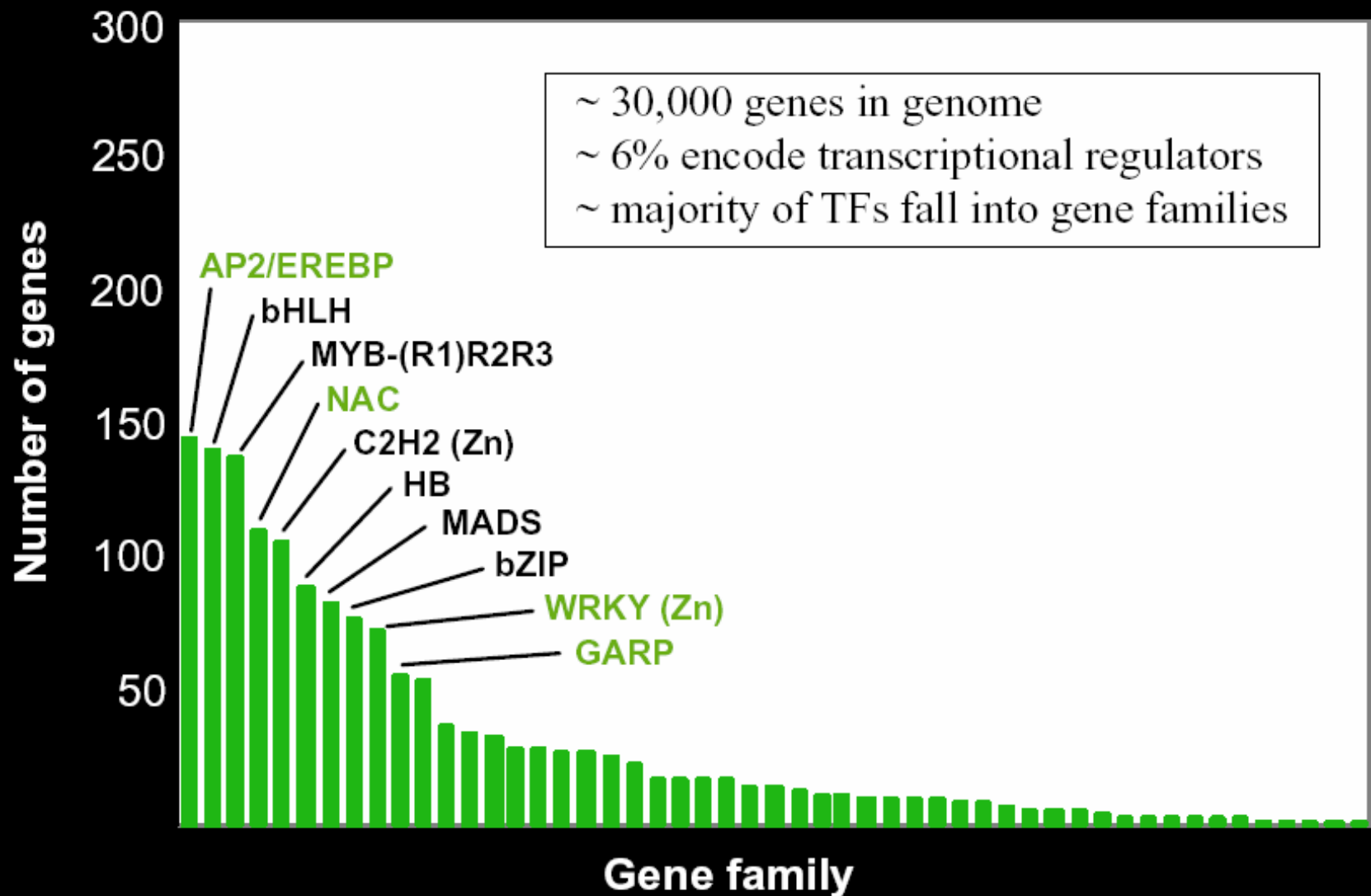
e.g. teosinte branched (CYC-family)
Doebley et al., 1997, Nature 386

Stress tolerance



e.g. CBF genes (AP2 family)
Stockinger et al., 1997, PNAS, 94

Arabidopsis thaliana transcription factor complement



Screens on transcription factors have produced hundreds of leads

Developmental Traits...

Branching pattern

OEX line wild type

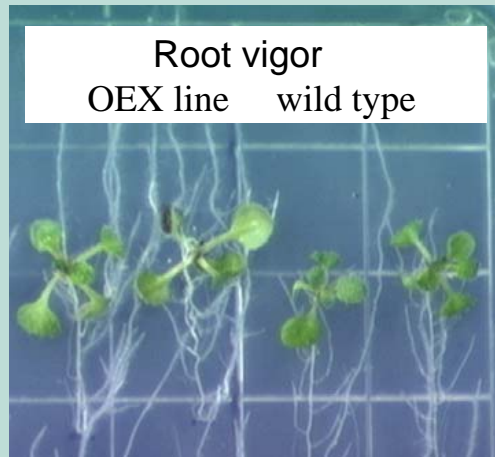
flowering time

OEX line wild type

seed size

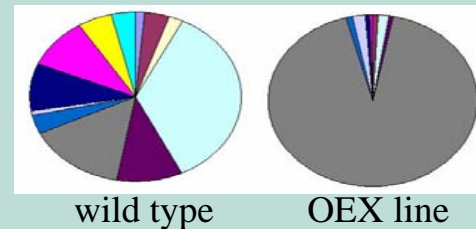


Root vigor
OEX line wild type



Biochemical Traits...

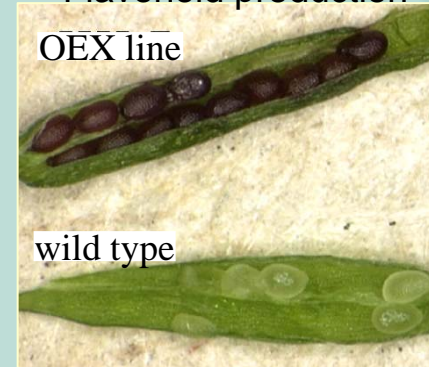
Glucosinolate composition



Flavonoid production

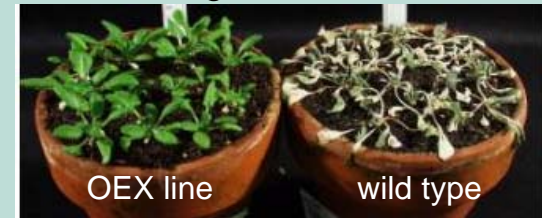
OEX line

wild type



Physiological Traits...

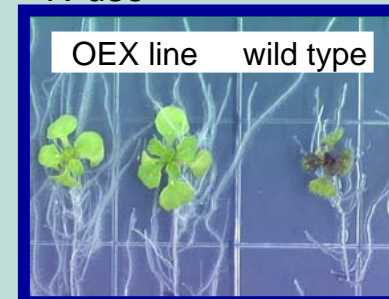
Drought tolerance



N-use

OEX line wild type

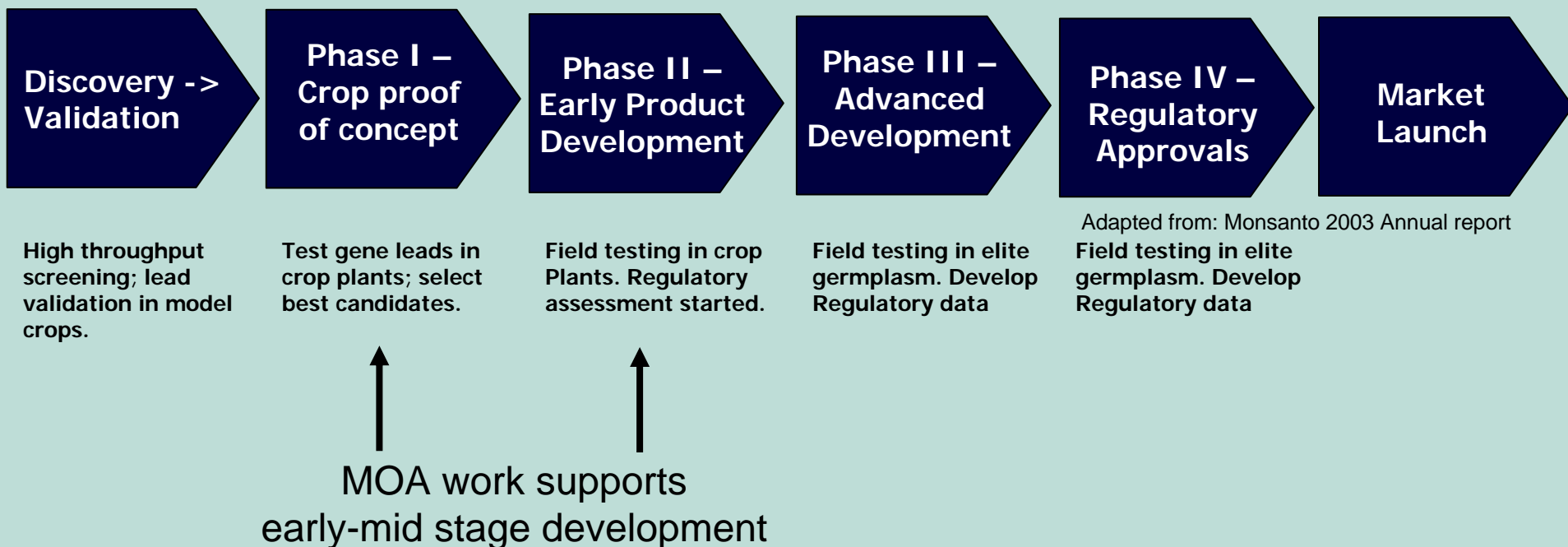
OEX line wild type



Typical Ag Biotech Product Development Process

Typical development timeline & costs:

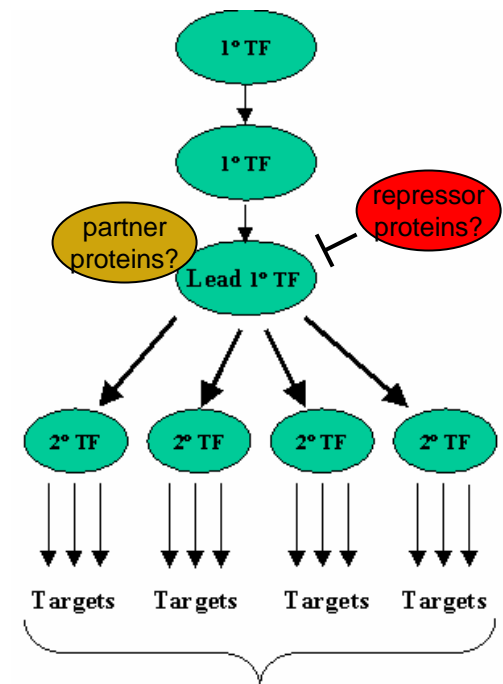
- 10 -15 years
- \$50M – \$100M



An understanding of MOA is important for successful commercialization

Mode of Action analysis:

- define phenotypic basis of the trait
- define pathways through which the leads act
 - => helps prioritize among leads
 - => support the future regulatory process/acceptance
 - => identify optimization strategies
- the genetic tools in Arabidopsis make it a valuable model



Trait of interest

Two general trait categories are of high priority

Intrinsic yield e.g.

- photosynthetic performance
- enhanced growth and vigor
- modification of architecture

Yield stability (stress tolerance)

-
- drought tolerance/water use efficiency
 - cold/freezing tolerance
 - N utilization
 - fungal disease tolerance

For recent review see:
Century et al., 2008
Plant Physiology 147, 20-29

We need a “Blue Revolution” (Kofi Annan, 2000)



- **Drought is the primary limitation to agriculture**
- **Many TFs have been implicated in drought responses**
 - Mostly from large TF families:
AP2/EREBPs, bZIP, NAC, MYB, MYC, C2H2, WRKY
 - Recent comprehensive reviews in:
Umezawa et al., 2006 and Bhatnagar-Mathur et al., 2008

NF-YB1 – a promising drought tolerance technology

Nuclear Factor Y (NF-Y) transcription factor family

- binds CCAAT box elements
- single copy TFs in yeast & mammals with roles in:
 - energy metabolism
 - cell cycle
- family greatly expanded in plants emerging roles in plant in:
 - flowering time
 - photosynthesis biology
 - drought responses

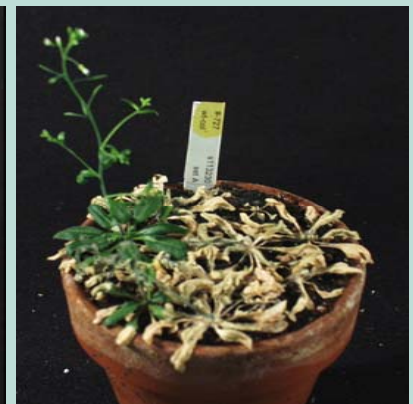
8 days
drought

35S::NF-YB1

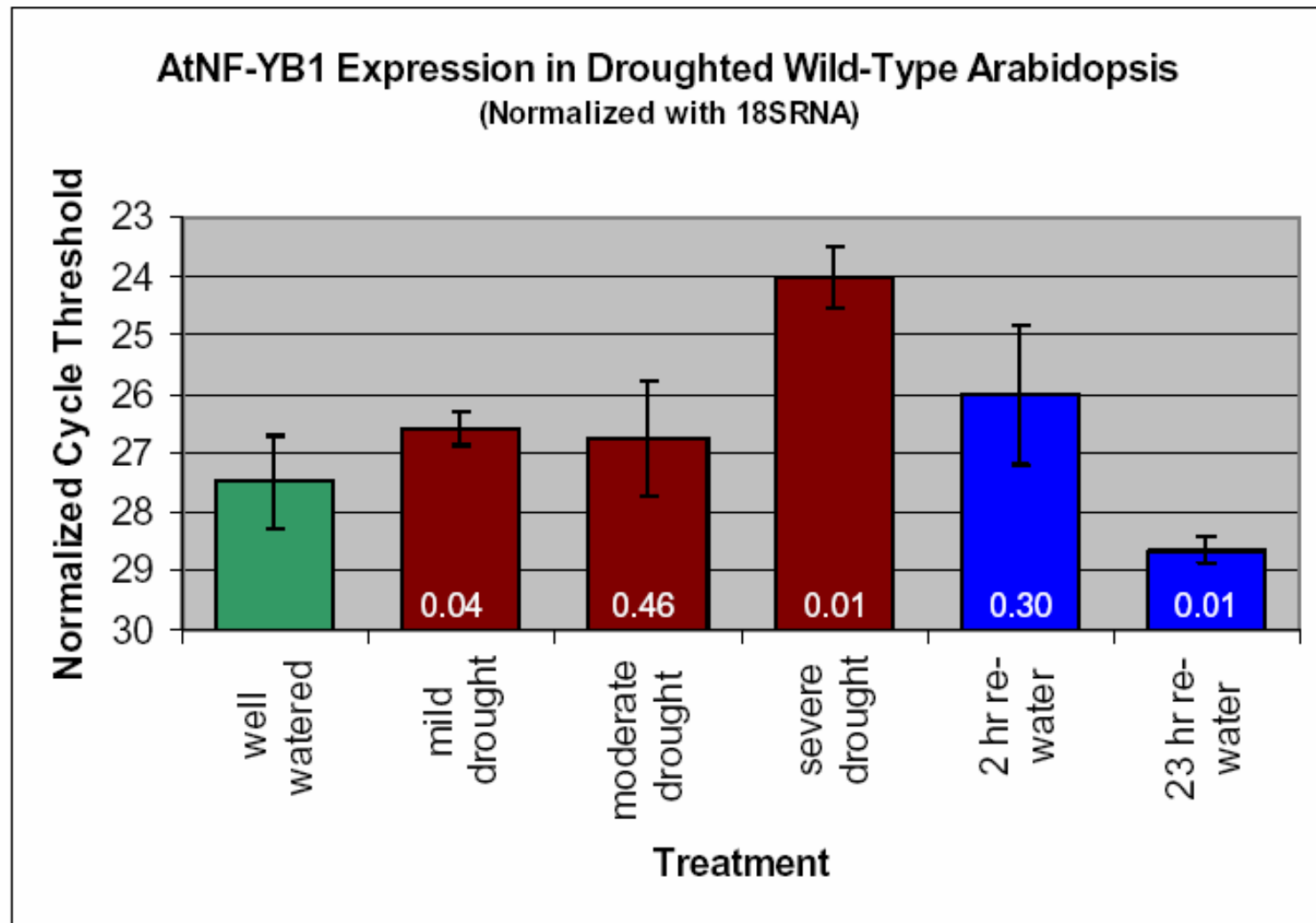
wild type



After
Re-water



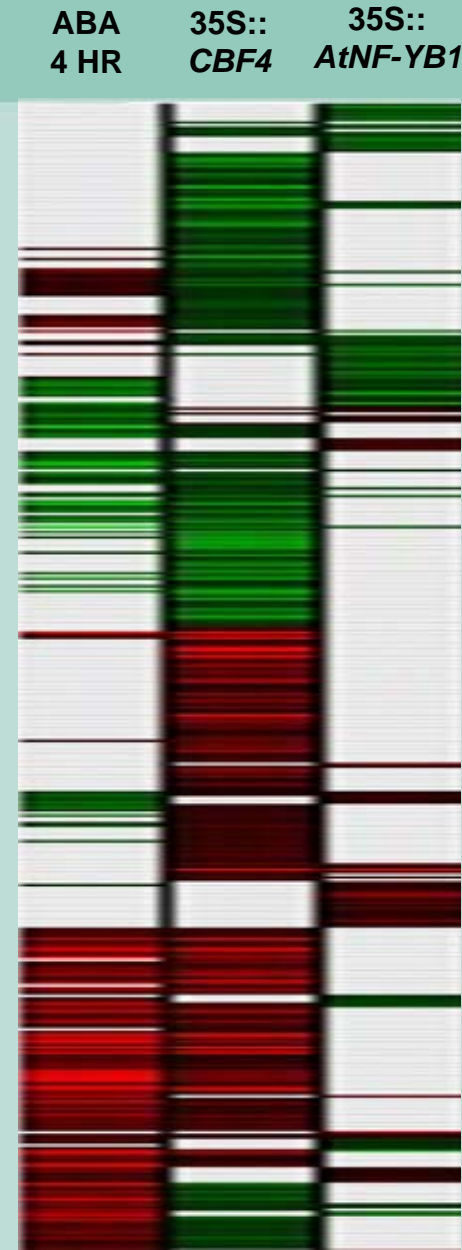
NF-YB1 is part of a native drought response



***NF-YB1* is upregulated ~12 fold by severe drought**

NF-YB1 regulates a novel genetic pathway for control of drought responses

- Microarray fingerprint shows little overlap with well characterized drought response pathways (ABA and CBF pathway).
- Key targets observed in 35S::*NF-YB1* profiles
 - photosynthesis components
 - pathways for synthesis of protective pigments
 - components of energy metabolism
- Key components of phenotype
 - increased WUE
 - increased chlorophyll levels
 - maintenance of turgor during stress
 - higher rate of photosynthesis during stress



Both The Genes and Control Pathways Are Conserved Across Arabidopsis....



Non-transgenic control

AtNF-YB1 Transgenic

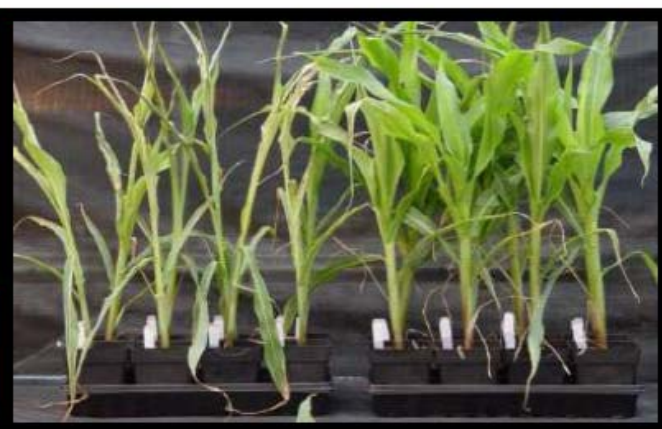
....soybean,



Non-transgenic control

AtNF-YB1 Transgenic

cotton...



Non-transgenic control

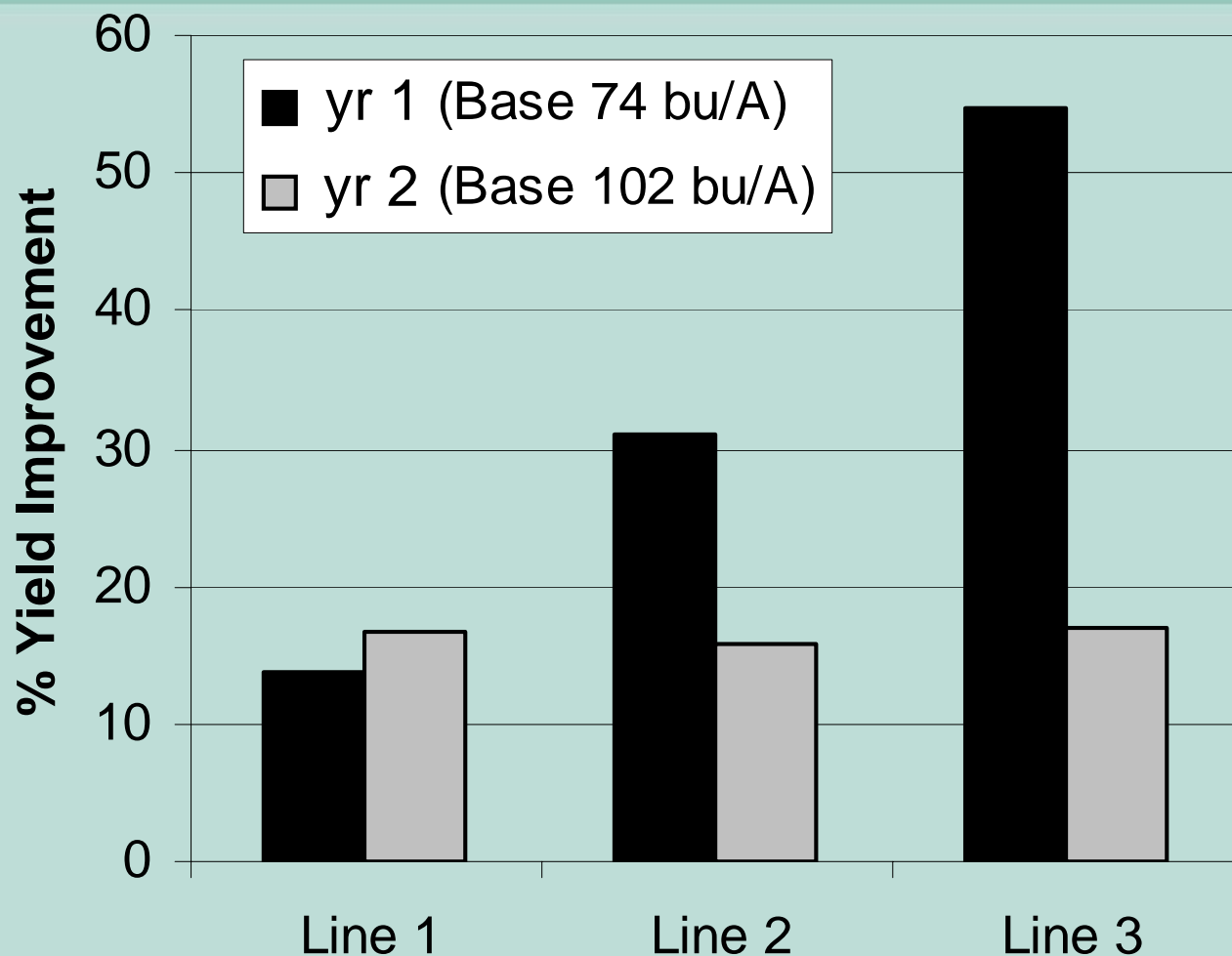
ZmNF-YB1 Transgenic

...and corn

Courtesy of Monsanto Company

Field trial results in corn

Significant Yield Advantage obtained on dry acres



Courtesy of
Monsanto Company

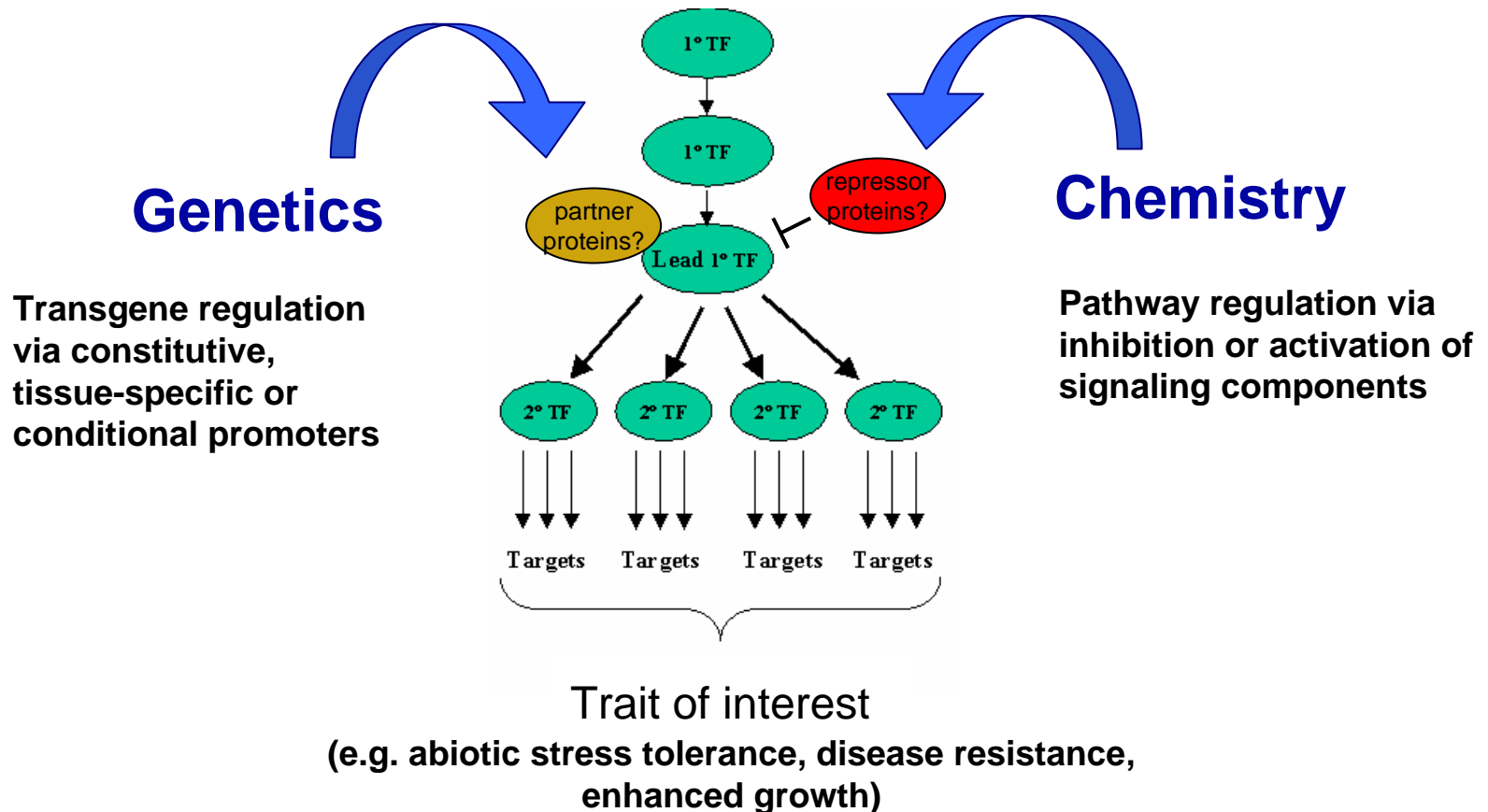
year 1: single, 6-replication drought efficacy trial using a Split-Split-Split block design in Kansas
year 2: multi-location study using group block design with 3 replicates within each location

New tools to meet the future goals of plant biotech

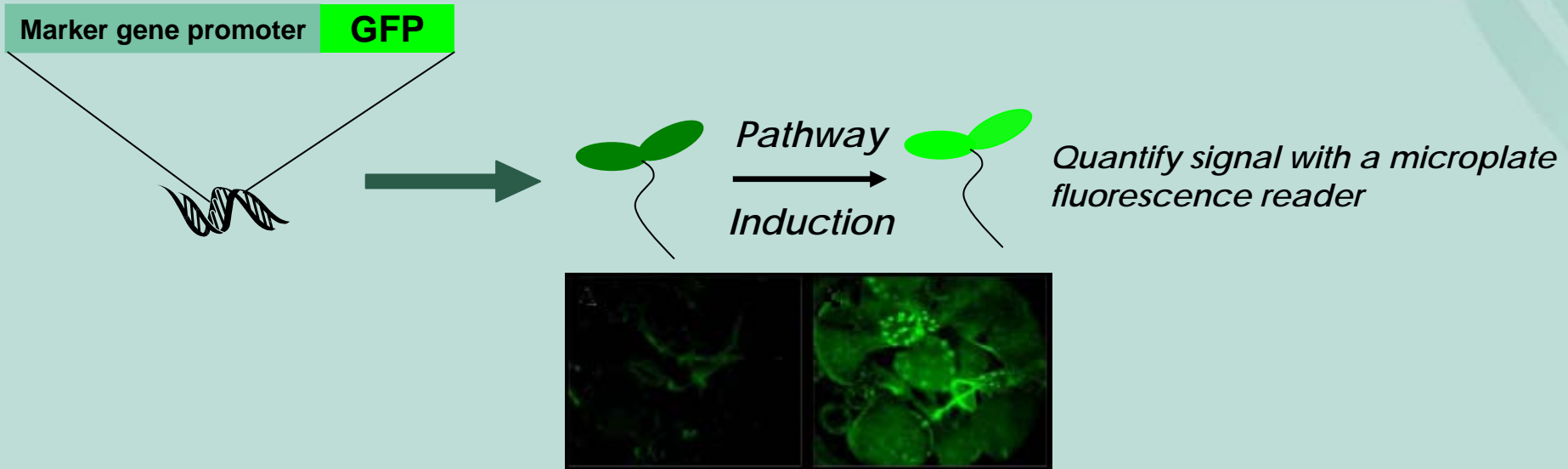
Excerpt from Monsanto press release from June 04, 2008

“ ... Monsanto will double yield in its three core crops of corn, soybeans and cotton by 2030, compared to a base year of 2000...”

Chemical approaches: complementary to genetics for modulating transcription factor (TF) regulated pathways



Reporter lines enable quantification of pathway induction



Promoters are selected using

- ~ Public data (genetic pathways)
- ~ Proprietary data (transcriptional profiling (TxP) analysis)

An ideal promoter

- ~ Conserved across species
- ~ Induced specifically in the pathway of interest
- ~ Expressed in leaf/cotyledon tissue
- ~ Downstream of multiple regulatory pathways

Compound imparts drought tolerance on soil-grown plants

Arabidopsis

DMSO



Compound (350 μ M)



Survival following drought treatment and rewater

Barley



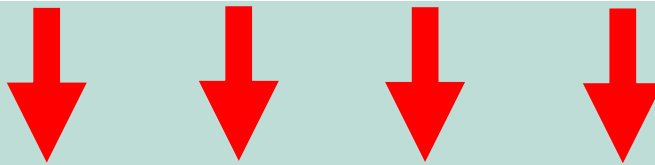
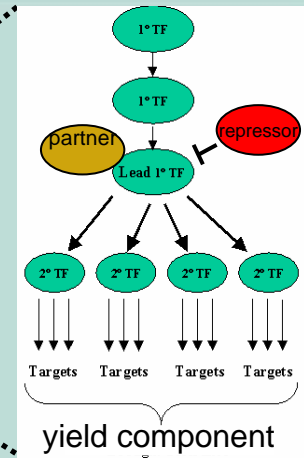
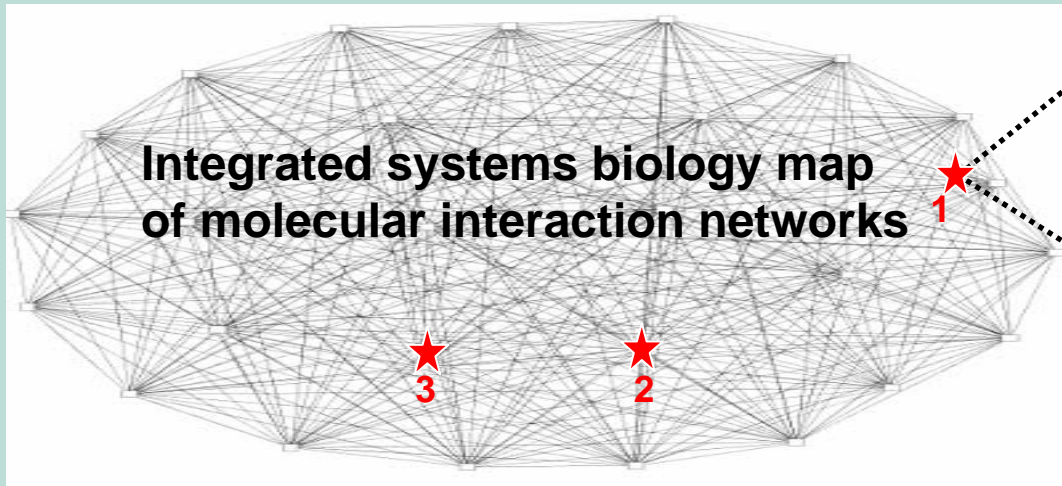
CONTROL

H144217
(500g/ha)

H144217
(1000g/ha)

Courtesy
Bayer
CropScience
AG

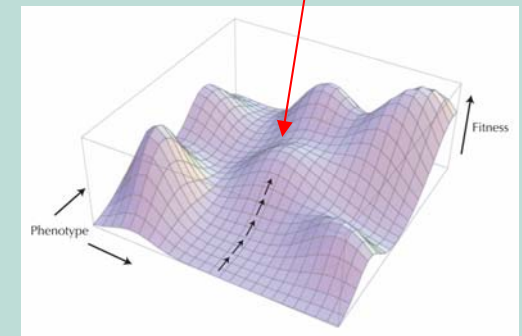
Systems biology will be important in delivering future technology



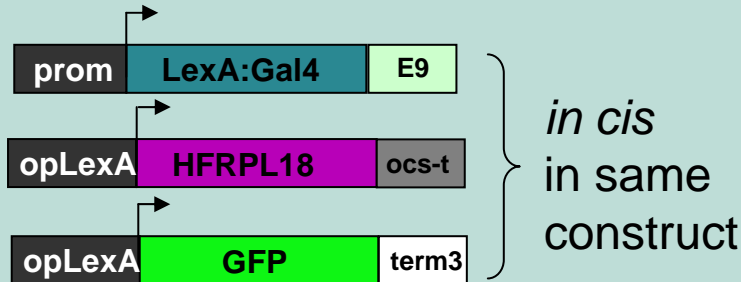
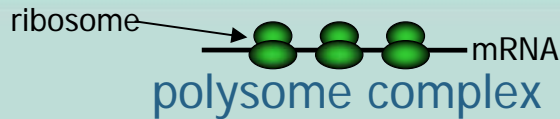
Network knowledge directs future rounds of discovery work

High resolution maps depend on high quality *in-vivo* TF x DNA and TF x protein interaction data at the **level of the individual cell-type**

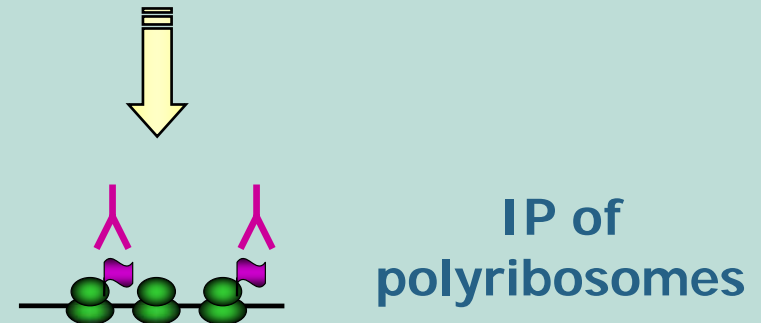
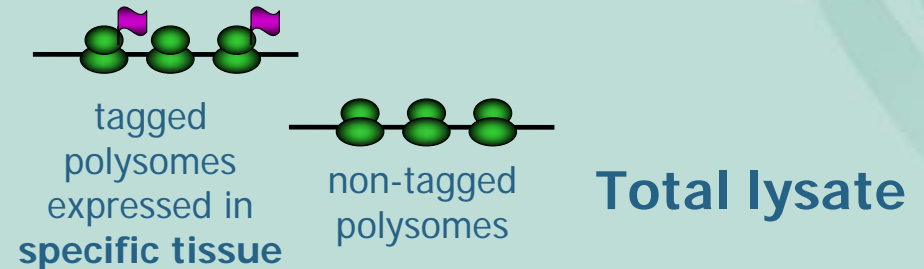
e.g. combination of
1 + 2 + 3 gives fitness peak



Ribotag: an emerging tool for cell-type specific expression studies



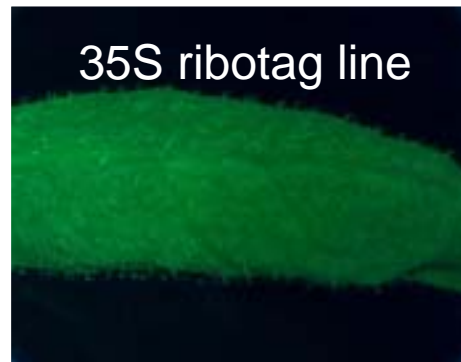
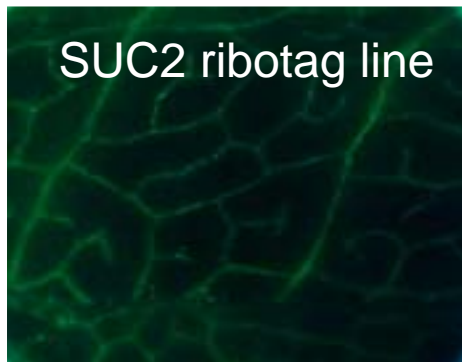
Stable transformation of
Arabidopsis



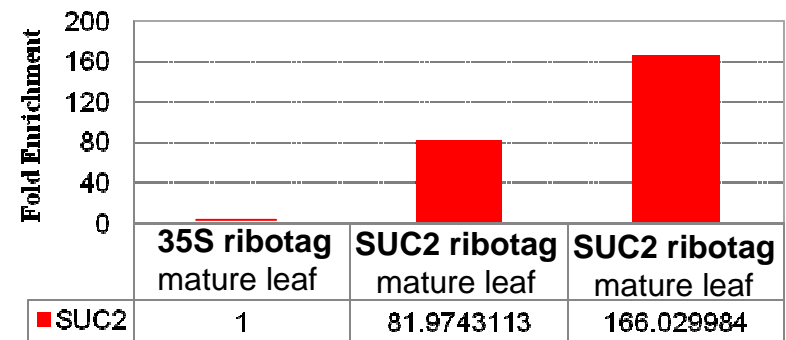
**RNA isolation and
expression analysis**

Reference: profiling of RNA associated with polysomes via tagging of RPL18
Zanetti et al. (2005). Plant Physiol 138, 624-635

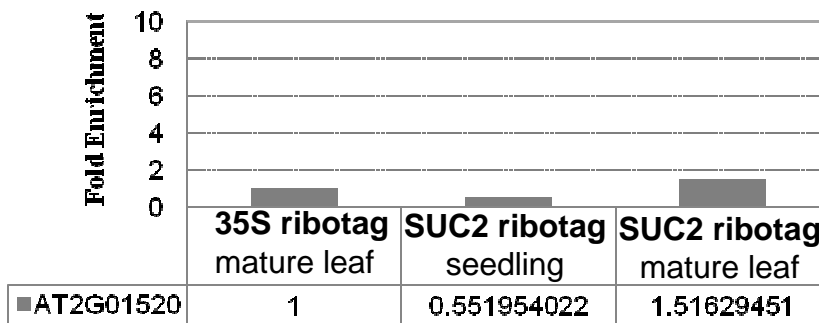
Validation: dramatic enrichment of specific markers in the phloem



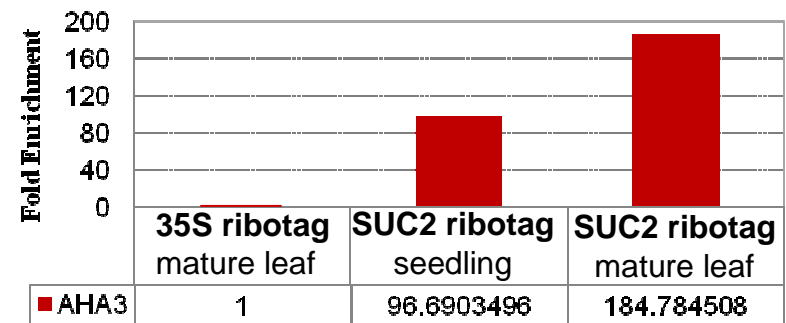
SUC2 Expression



ATG01520 Expression



AHA3 Expression



Projected timelines for biotechnology products

Impact
of
molecular
biology



first generation (mid 1990s)

- **single gene traits – herbicide and insect resistance**

Impact
of
Genomics



next generation (early 2010s)

- **complex traits – yield and stress**
- initial lead technologies identified from screens

Impact
of
Systems
Biology
&
Chemistry



later generations (> 10 years hence)

- **super-enhanced complex traits – yield and stress**
- optimal intervention points in regulatory pathways
- rationalized from “holistic” knowledge of entirety of pathways

Concluding remarks



- TF-based technologies are excellent candidates for enhancement of crop yield and stress tolerance
- Genetics and chemistry offer complementary approaches to pathway regulation
- Systems biology will enable future generations of super-enhanced products
- Plant biologists have a critical role to play in ensuring a sustainable future**