



INNO-FOOD

Plant Breeding

Madesis Panagiotis
Researcher





Plant Breeding

- Conventional breeding
- Molecular Breeding
- Genetic Engineering





What is plant breeding?

- Genetic improvement through crossing plants with desired traits and selecting progeny with improved performance and/or improved combinations of traits.
- “Accelerated” and “targeted evolution”.
- Application of genetics principles to crop improvement.
- Systematic procedures used to improve trait phenotypes by crossing and selection, directed manipulation of the genotype at the DNA sequence level, and introduction of new genes.





Do we still need to train students in
plant breeding?





Agricultural food production

- Each year humans re-create the food supply that feeds 6.15 billion people
- Reserves of staple foods would feed the world for less than two months and were as low as 48 days in 1995
- 800 million people go to bed hungry every night

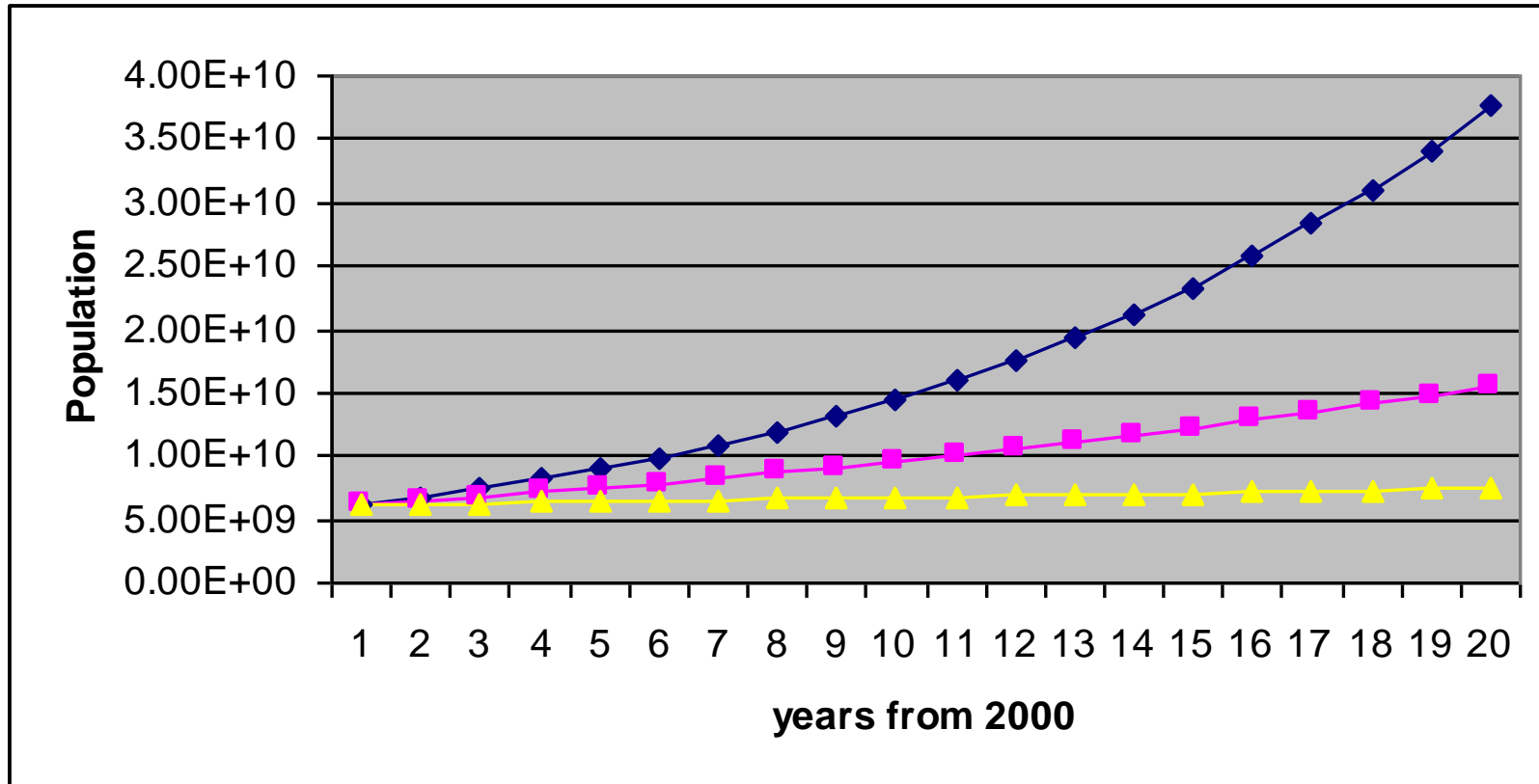
Food and Human nutrition

- <http://www.harvestplus.org/>
- Vitamin A deficiency affects 750 million people





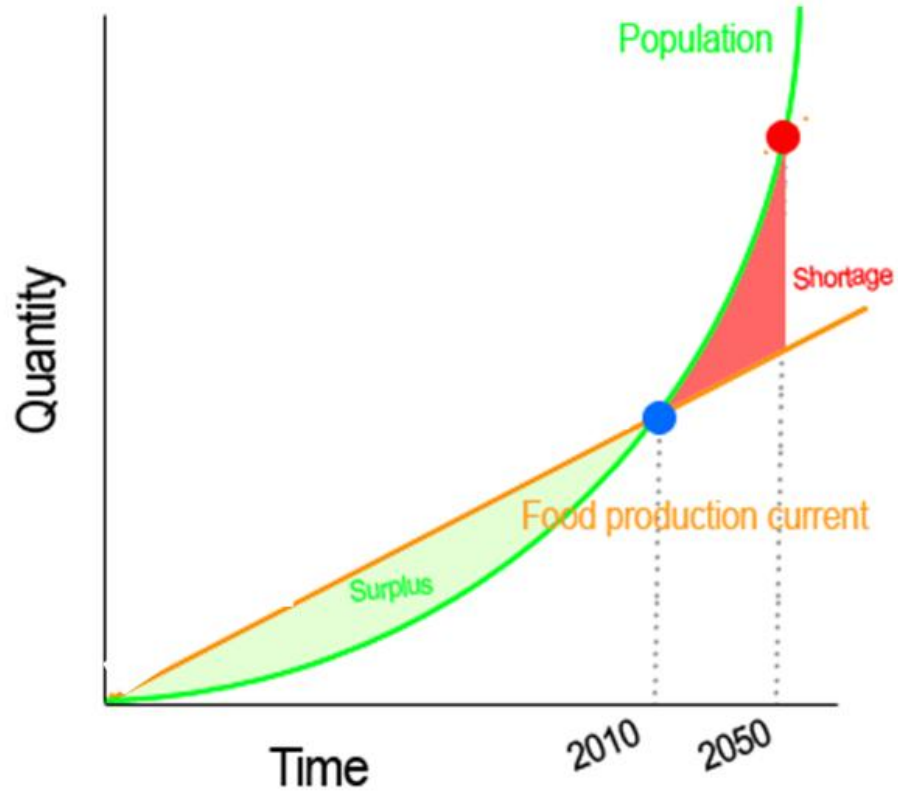
Projected population increase under different assumptions of reproduction and mortality





Food
Τρόφιμα

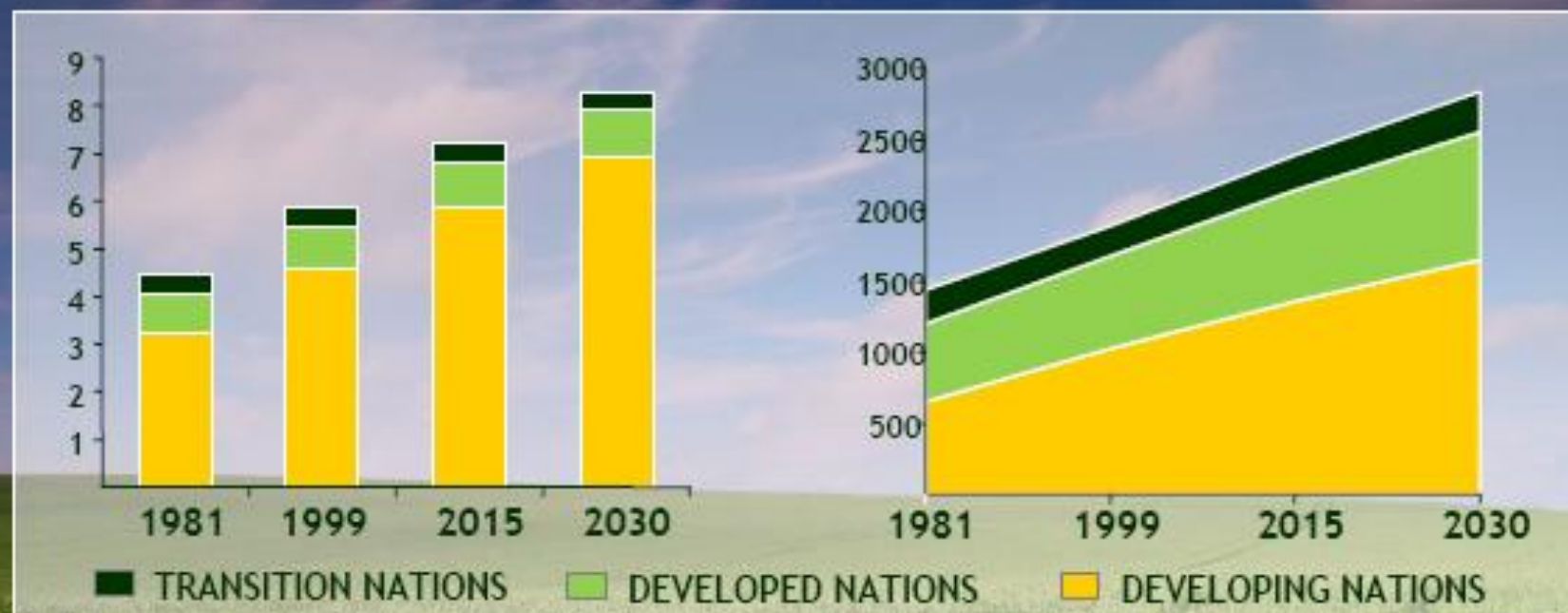
Current Food Production vs. Future Food Production In Relation to Growing World Populations Over Time



In the next 50 years we have to produce more food than ever before in the history of humankind

GROWING WORLD POPULATION (B)

RIISING CEREAL DEMAND (MMT)



- **World population continues to increase**
- **Per capita food consumption continues to rise**
- **Consumers continue to demand improved taste, convenience, and nutrition**

Source: FAO, WHO



Agricultural F production



Food
Τρόφιμα



Feed
Τροφές



Fiber
Ίνες



Flowers
Άνθη



Farmaceuticals
and Parfums
Φάρμακα και Αρώματα



Fuels
Καύσιμα





Projections

- We need to make as much progress in production efficiency in the next 30 years as we have made in the previous 12,000
- We need to double food production by 2050





The success of plant breeding

- Increases in yield are derived both from improved varieties and from improved management.
- In vegetable crops, research suggests about a 50-50 split between genetic gain and gain attributed to management.
- Genetic gain in grain yield of $75 \text{ Kg ha}^{-1} \text{ yr}^{-1}$ for corn can be attributed to breeding.
 - 1 ton/acre increase in yield every 30 years.
 - Maize yields have increased 60% to 120% in the U.S. since 1940 (Cooper et al., 2004. Genomics, Genetics, and Plant Breeding: A Private Sector Perspective. Crop Science. 44:1907-1913).
- “Green revolution” varieties have increased yields 2 to 3 fold in many “developing” nations (Knight, 2003).





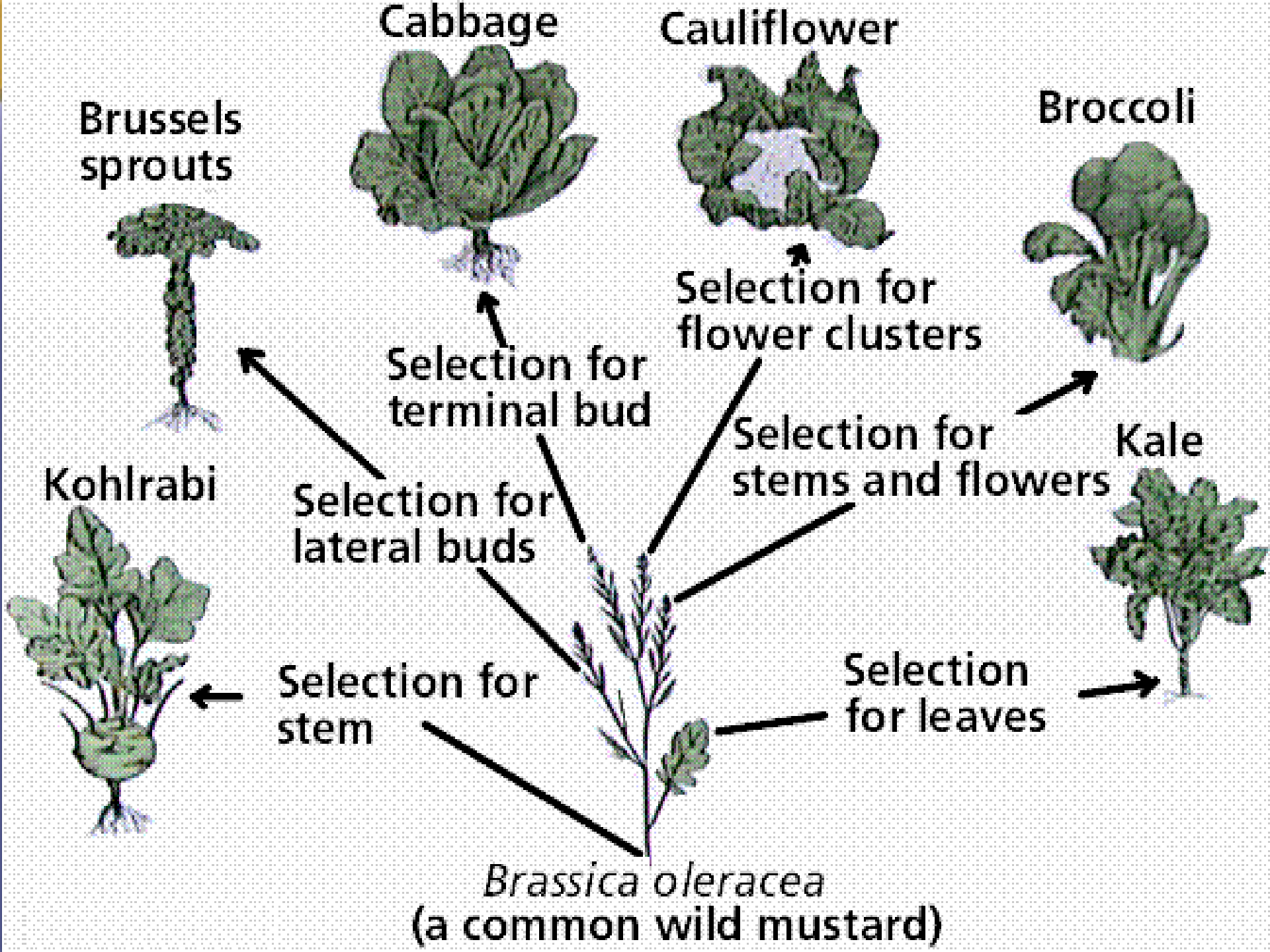
Green revolution



Norman Borlaug

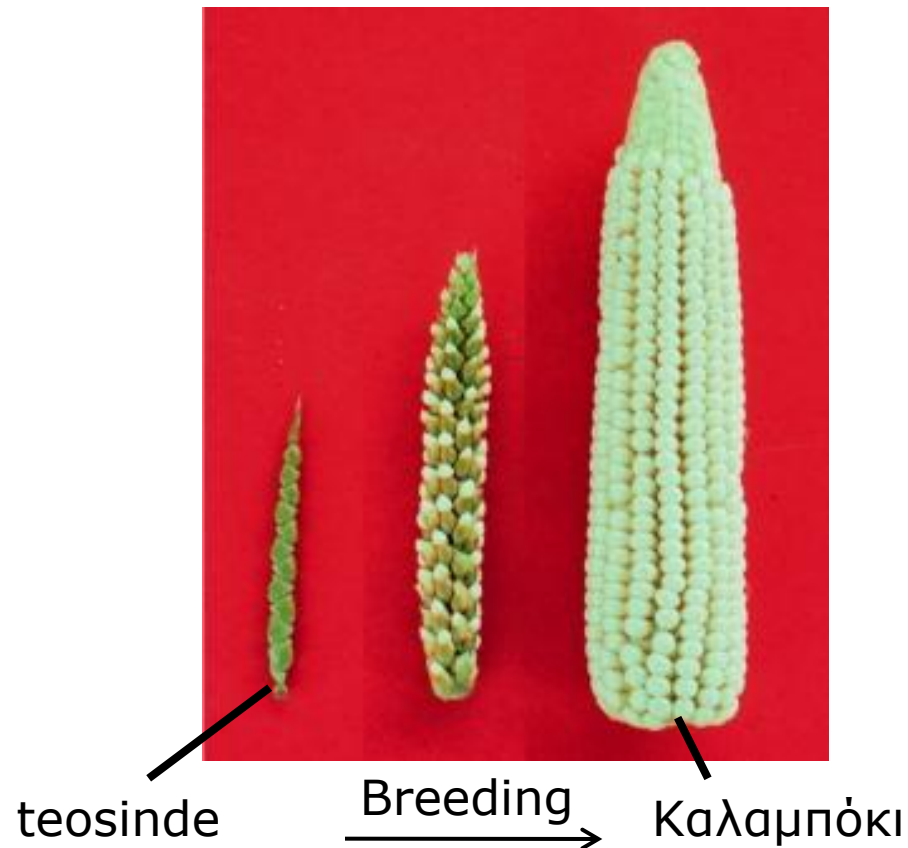
“Short” wheat disease resistant ('50s)







Breeding success story





Breeding success story

wheat



Aegilops tauschii



Einkorn
AA

Emmer
AABB

spelt
AABBDD



Triticum aestivum
AABBDD





Is conventional breeding obsolete?

- Easy traits to manipulate via GMO techniques are single gene – these are also easily manipulated using conventional breeding.
- Conventional breeding can manipulate multiple traits simultaneously
- Conventional breeding can manipulate genetically complex “quantitative traits”

Traits that are influenced by the environment

Traits that are conditioned by multiple genes

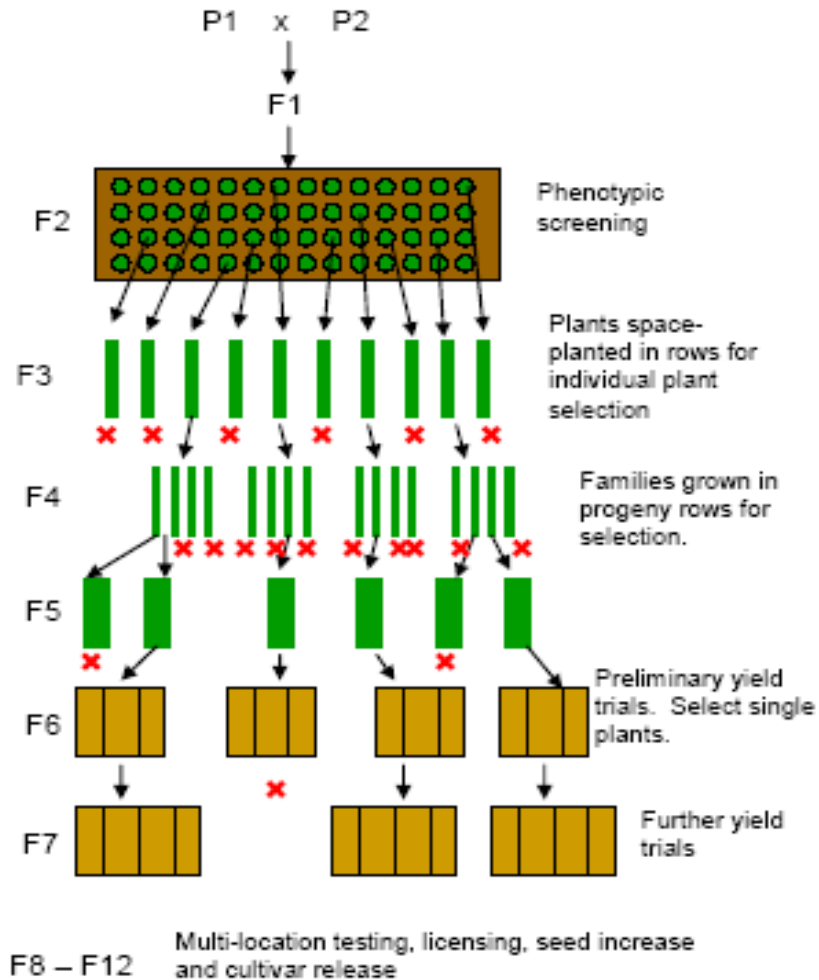
- Selection on phenotype is a powerful approach to bring about directed changes. (Robust but can be slow; requires that genetic variation exist for the trait of interest)
- Complex genotype x environment systems that agriculture operates under means that “methodology” of evaluation will always be important.



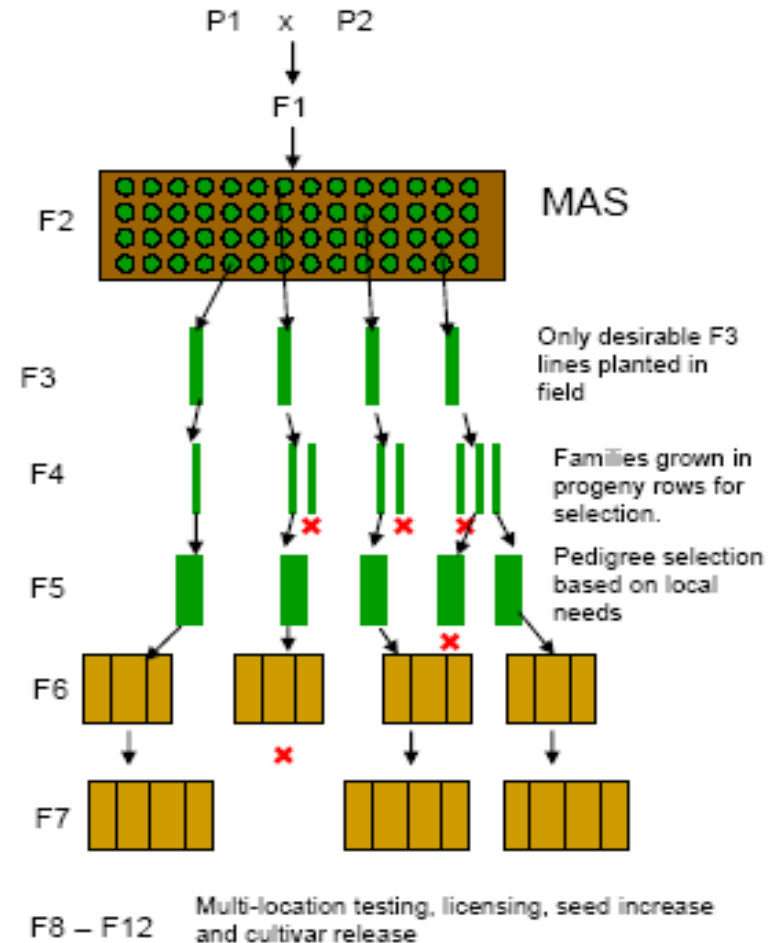


Early generation marker assisted selection

PEDIGREE METHOD



EARLY GENERATION SELECTION MARKER ASSISTED SELECTION



Early generation selection scheme (proposed by Ribaut & Betran (1999)). Note that many lines can be discarded in an early generation which permits the evaluation of fewer lines in later generations.



Molecular Breeding

What is a microsatellite?

- **Repeats with a motive (1-6 repeats)**

Dinucleotide	(CT) 6	- CTCTCTCTCTCT
Trinucleotide	(CTG) 4	- CTGCTGCTGCTG
Tetranucleotide	(ACTC) 4	- ACTCACTCACTCACTC

- **Classes**

Perfect repeat when repeat tract pure for one motif

CTCTCTCTCTCT

Compound SSR when repeat tract pure for two motifs

CTCTCTCACACA

Imperfect SSR if single base substitution

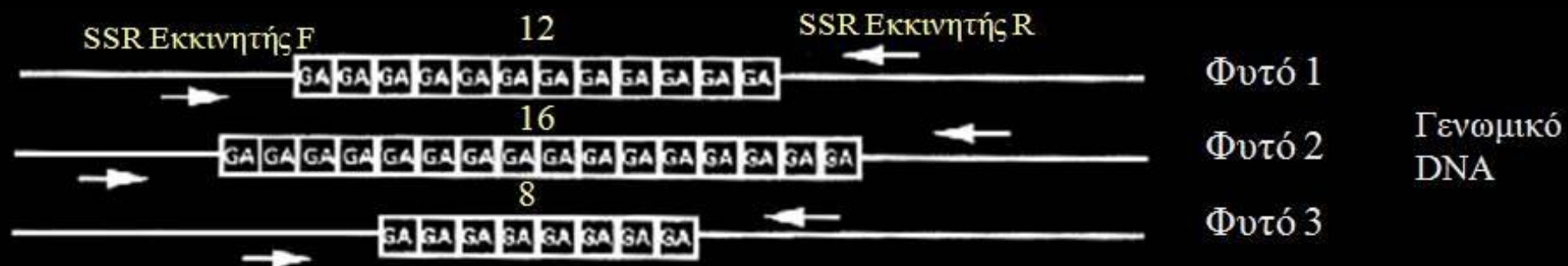
CTCTCTACTCTCT

Region of cryptic simplicity if complex but repetitive structure

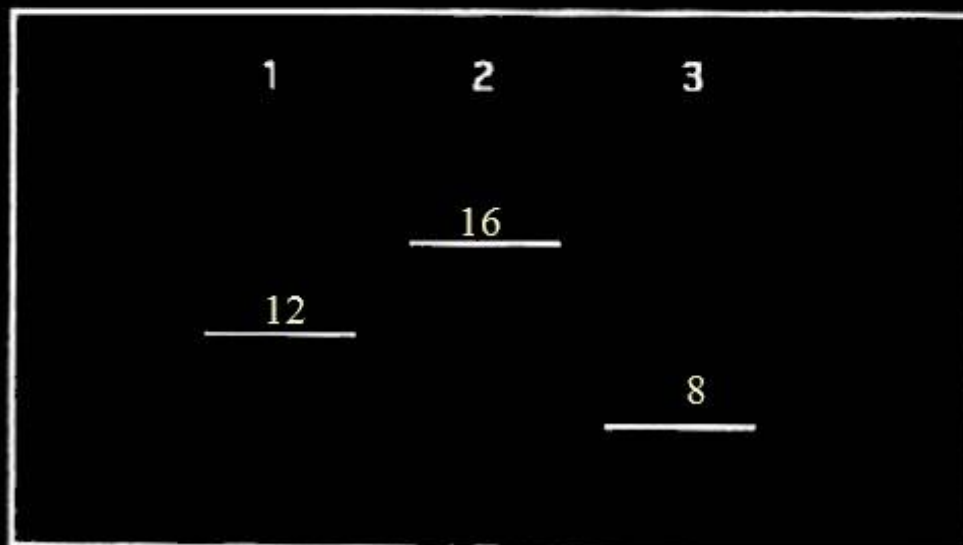
GTGTCACAGAGT



SSRs



Ενίσχυση μέσω τεχνικής της Αλυσιδωτής Αντίδρασης της Πολυμεράσης (PCR)



Πηκτή ανάλυσης

Metaphor Αγαρόζη

Πηκτή Πολυακρυλαμίδης



Ανάλυση με πρόγραμμα (software)

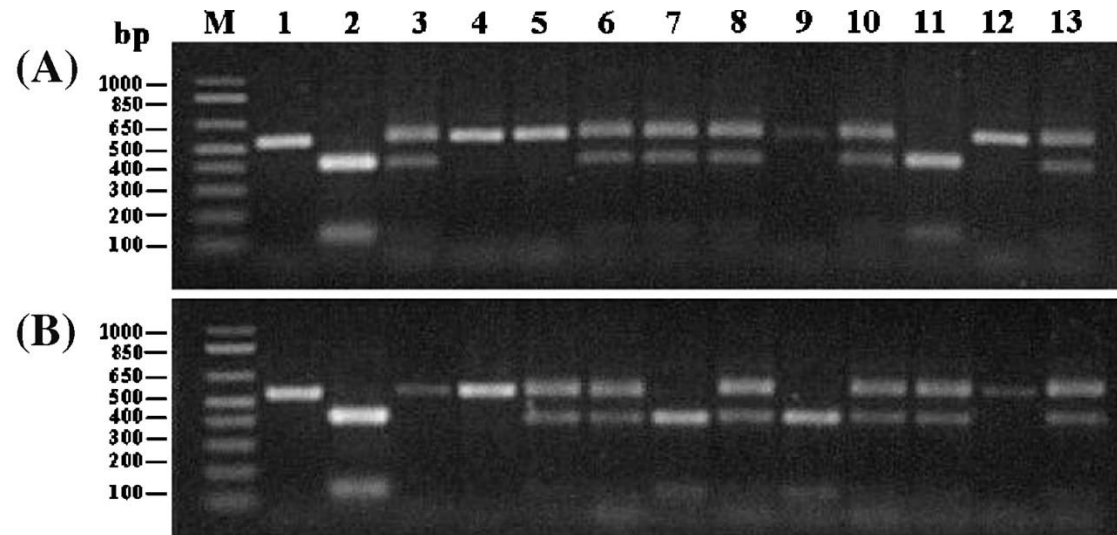


Εύρεση αριθμού και μεγέθους διαφορετικών Αλληλομόρφων



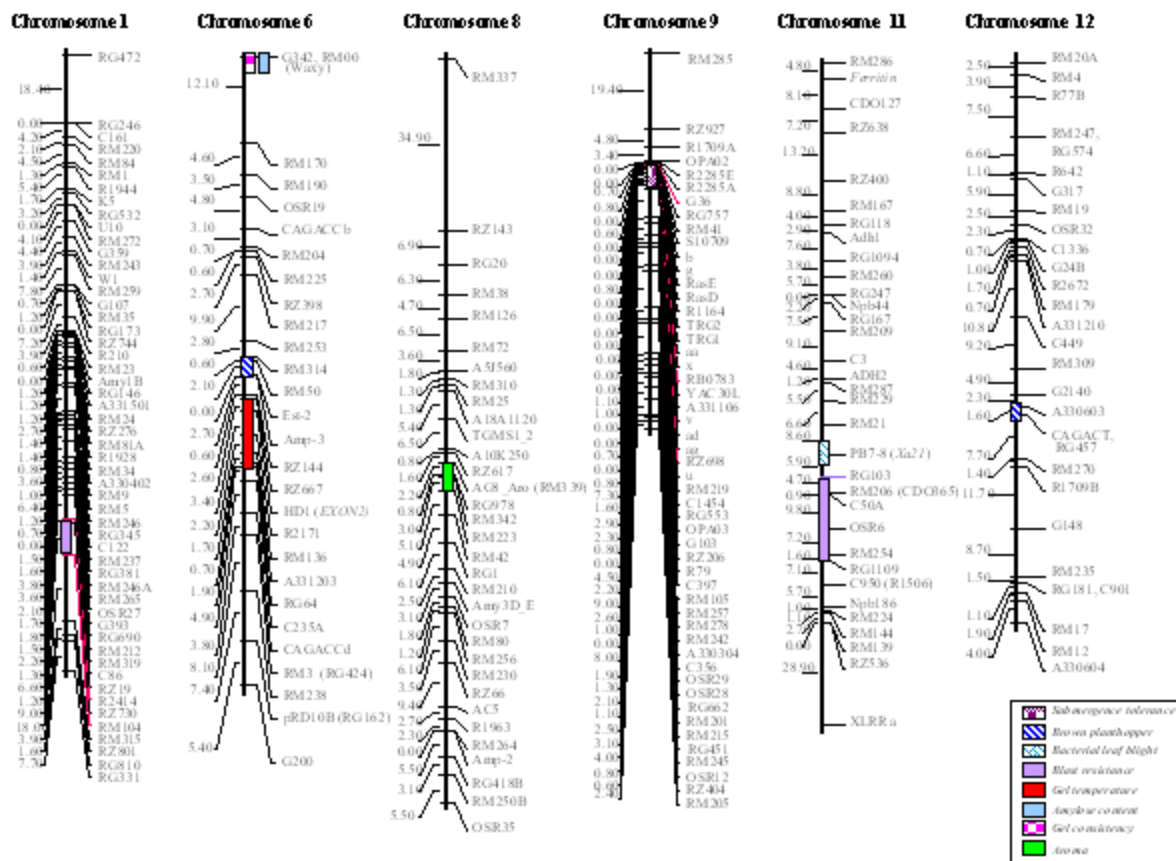
Molecular breeding

Marker assisted selection





Molecular Breeding

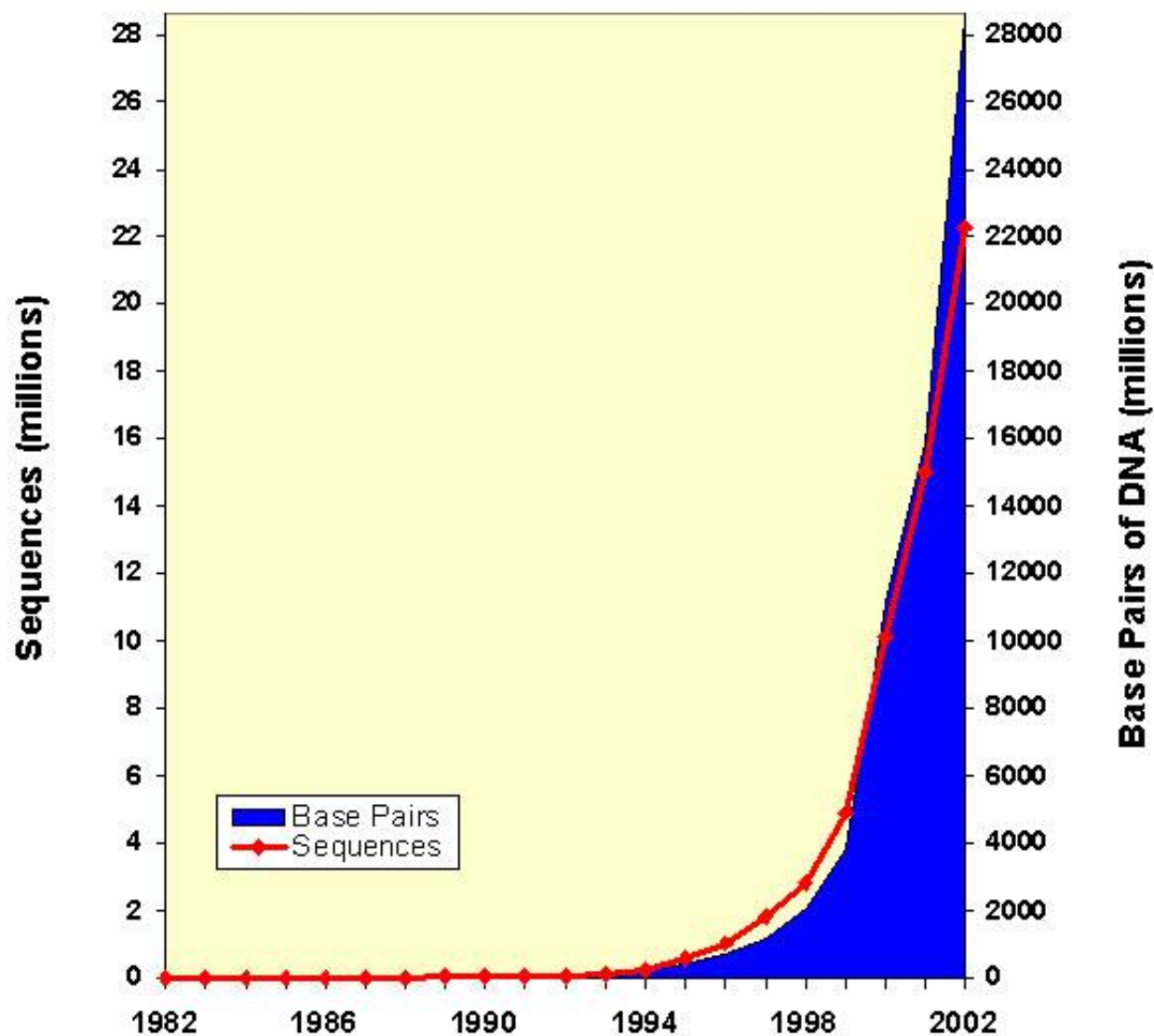


Using genomic maps



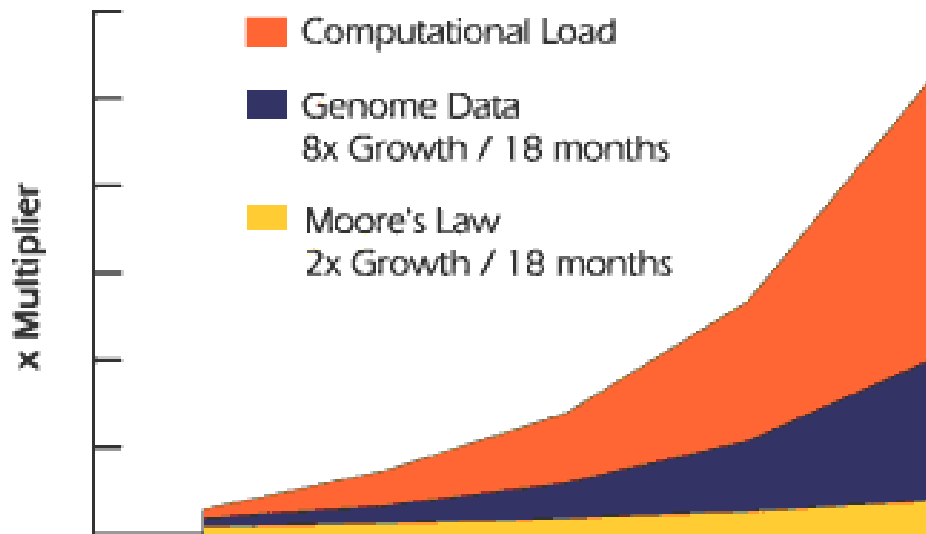


Growth of GenBank





The challenge of scale - computational

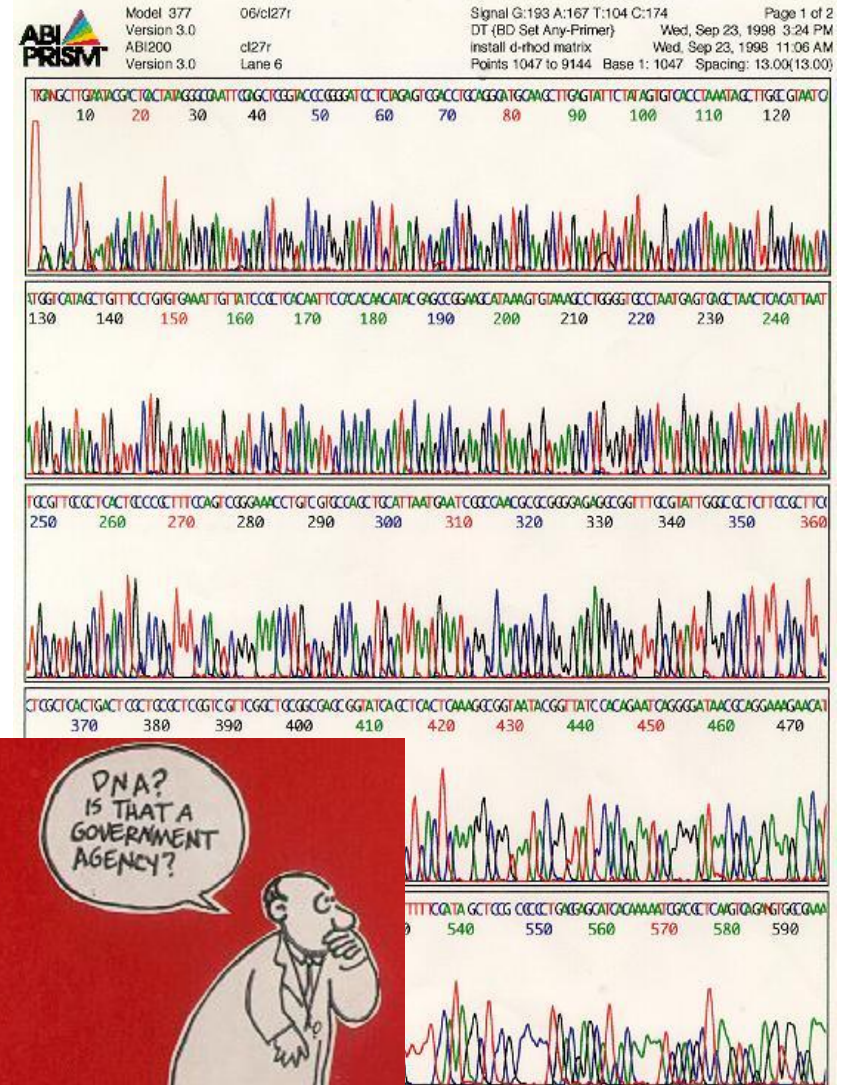


- High-throughput
- Pipelines
- Standardization
 - Controlled vocabulary
- Validation



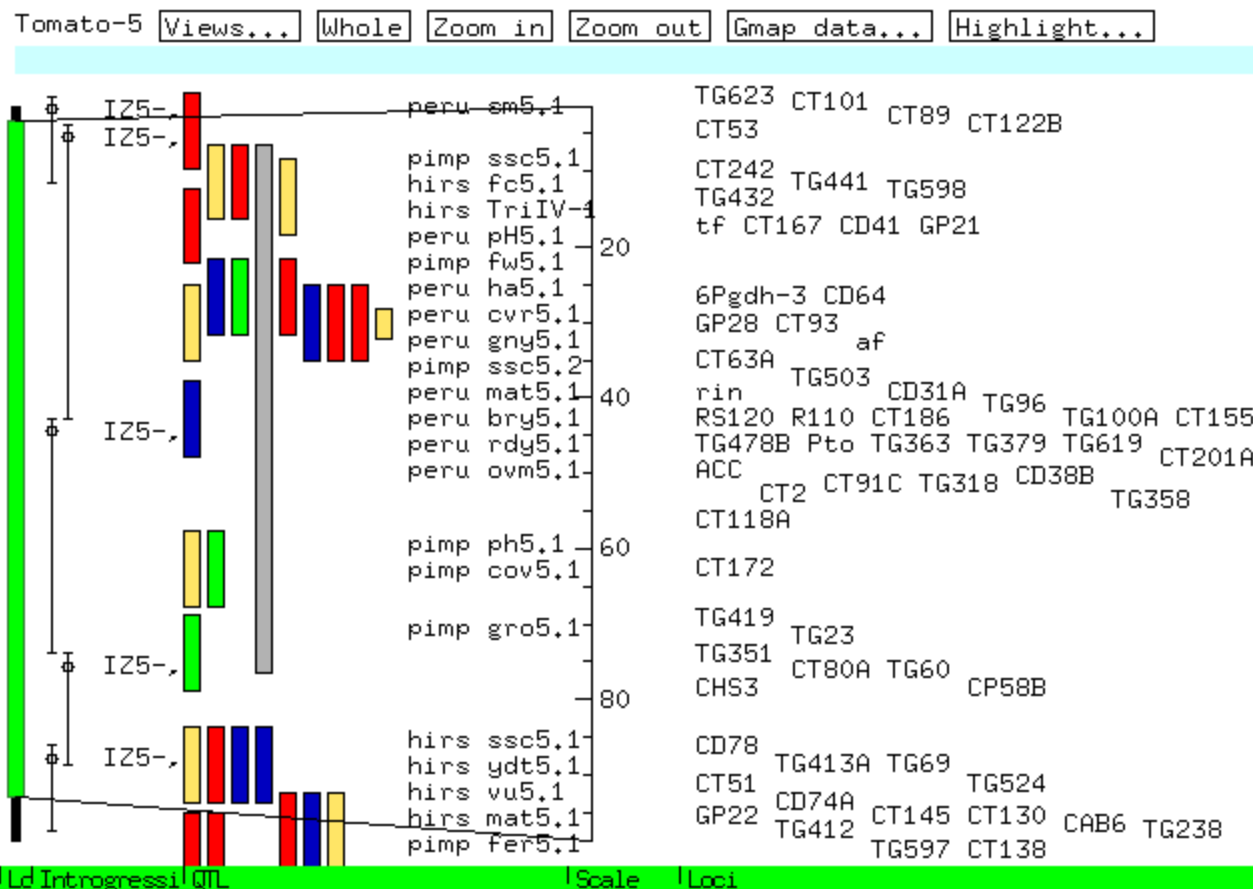


How can we use new information ?



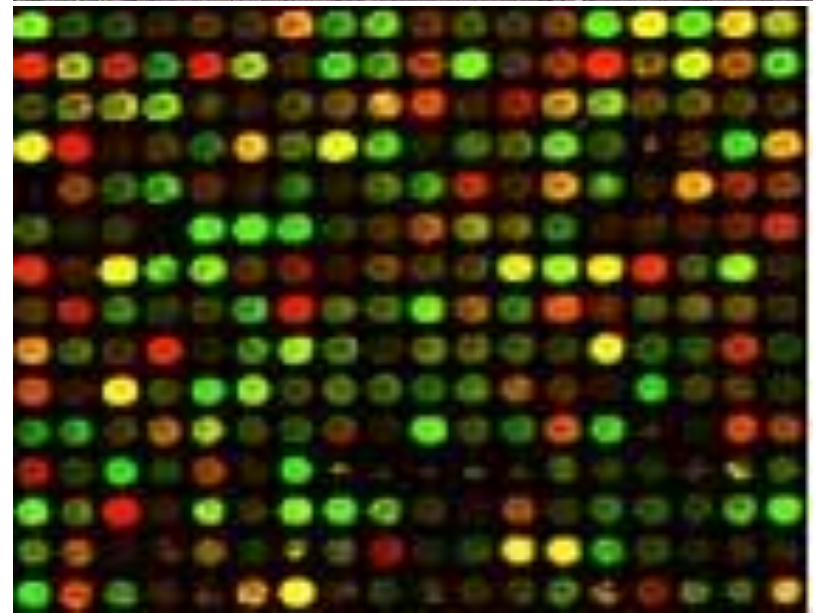


The results of “Structural Genomics”
gives us many new tools to improve
crops through “map-based” breeding



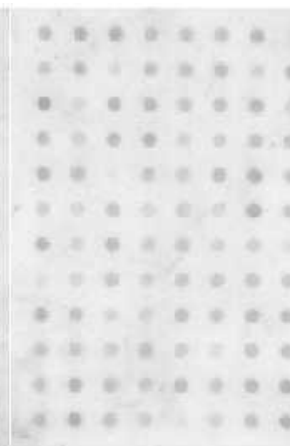
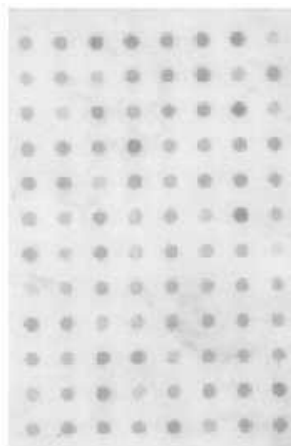


There is a broader context to the methodology e.g., quantitative methods used in plant breeding are currently being applied to the analysis of “DNA chip” experiments





- “When considering the handling of undesirable variation it is inevitable that the discussion will centre on agricultural field trials, since modern experimental technique was initiated and has reached its greatest elaboration in this realm”
- K. Mather, The Control of Error in Statistical Analysis in Biology, University Paperbacks, Methuen & Co., LTD., London.





Tobacco plant breeding

Combination

conventional

Marker assisted selection

Study genetic diversity

Parent selection

Parent selection

Crosses

Assessment of parents and progeny

Selection

back crosses

selection

Assessment of parents and progeny

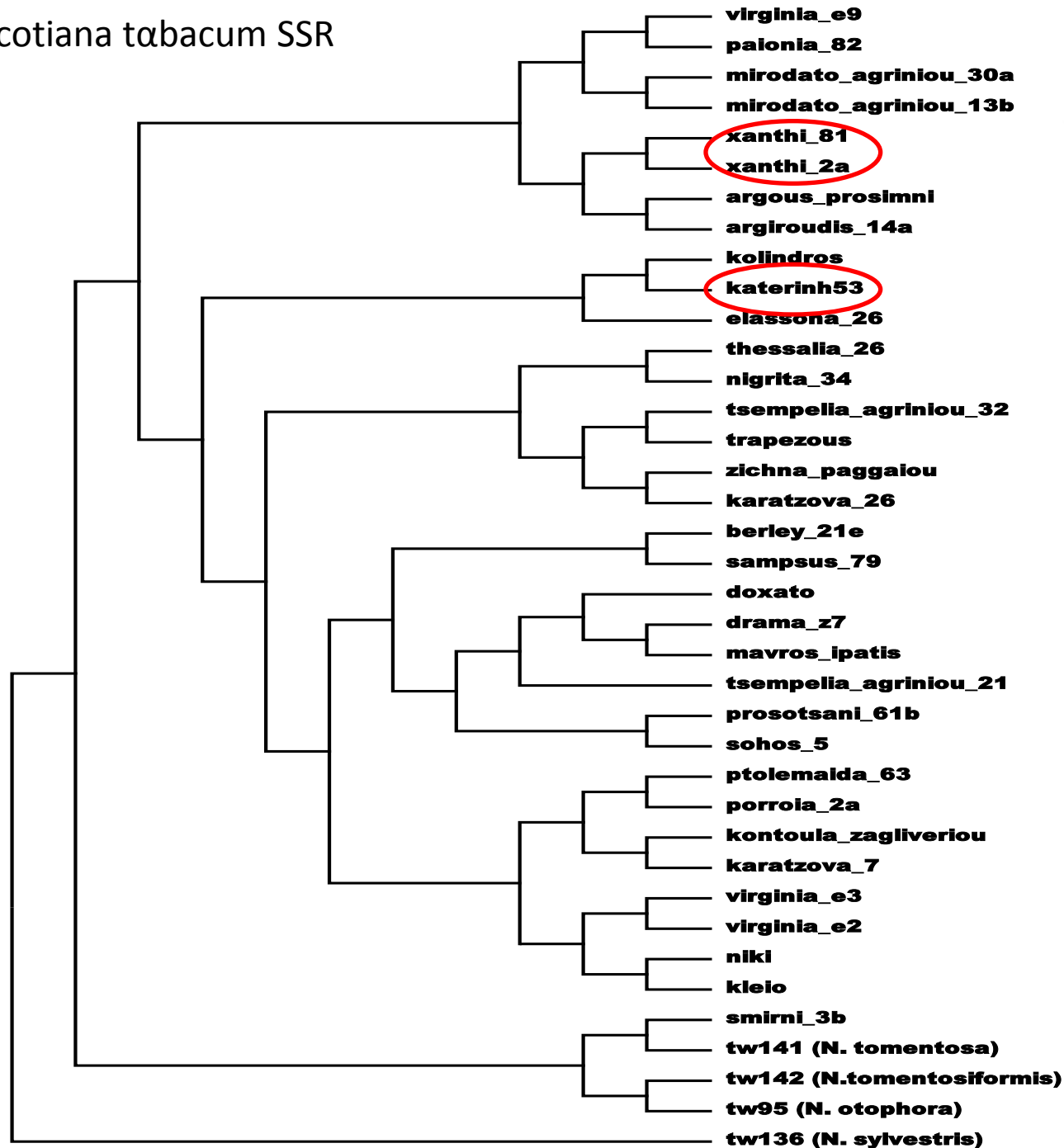
Pure lines

Assessment of progeny



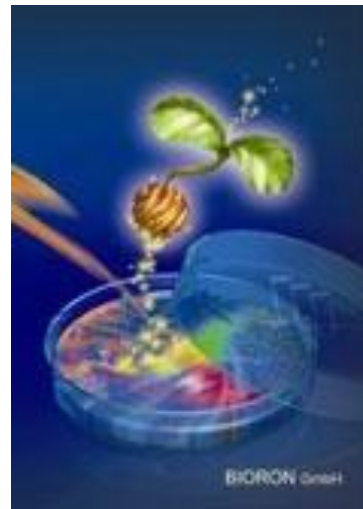


Nicotiana tabacum SSR





Genetic Engineering





Developing disease resistant plants

- Over expression of cpd1 as a means for cercospora resistance





Cercospora infection



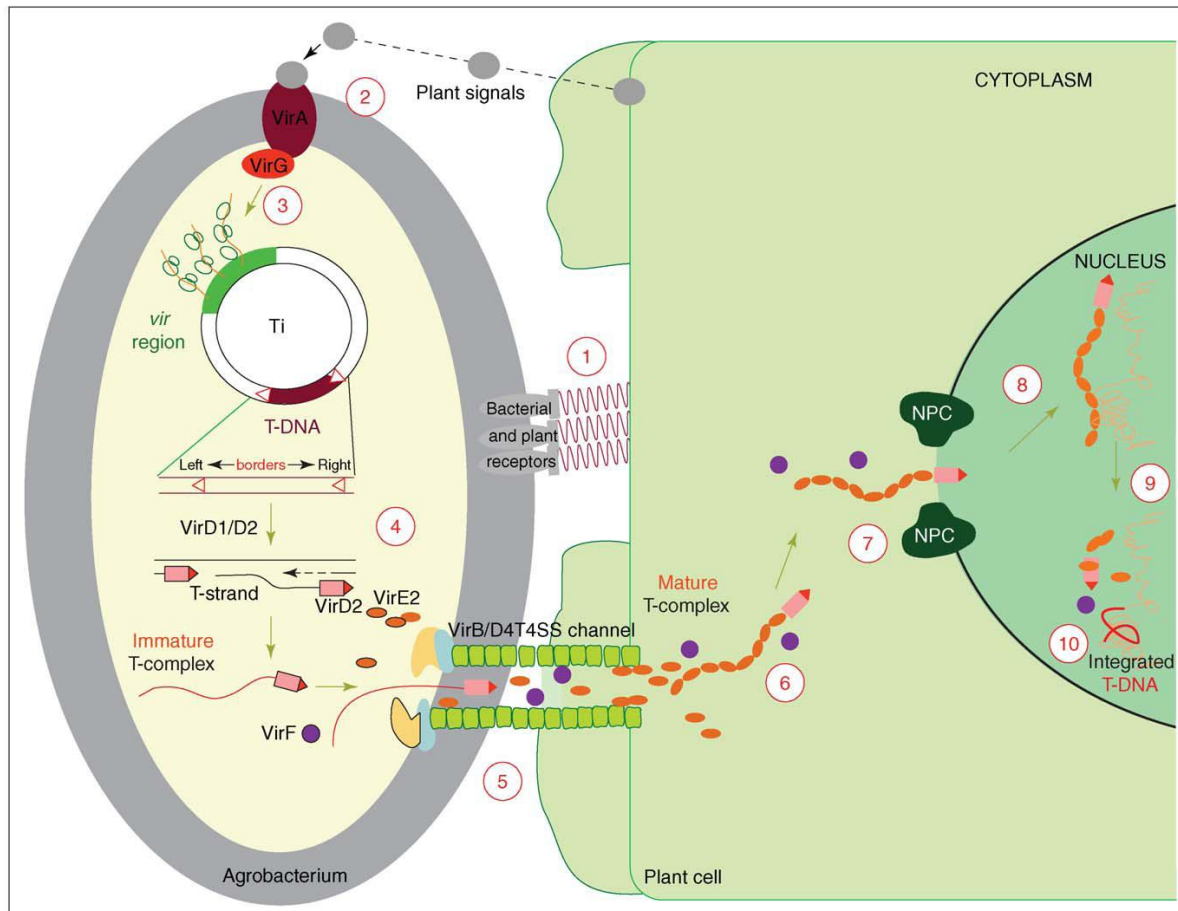


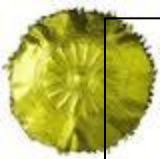
Cercopsora on beta vulgaris leaves



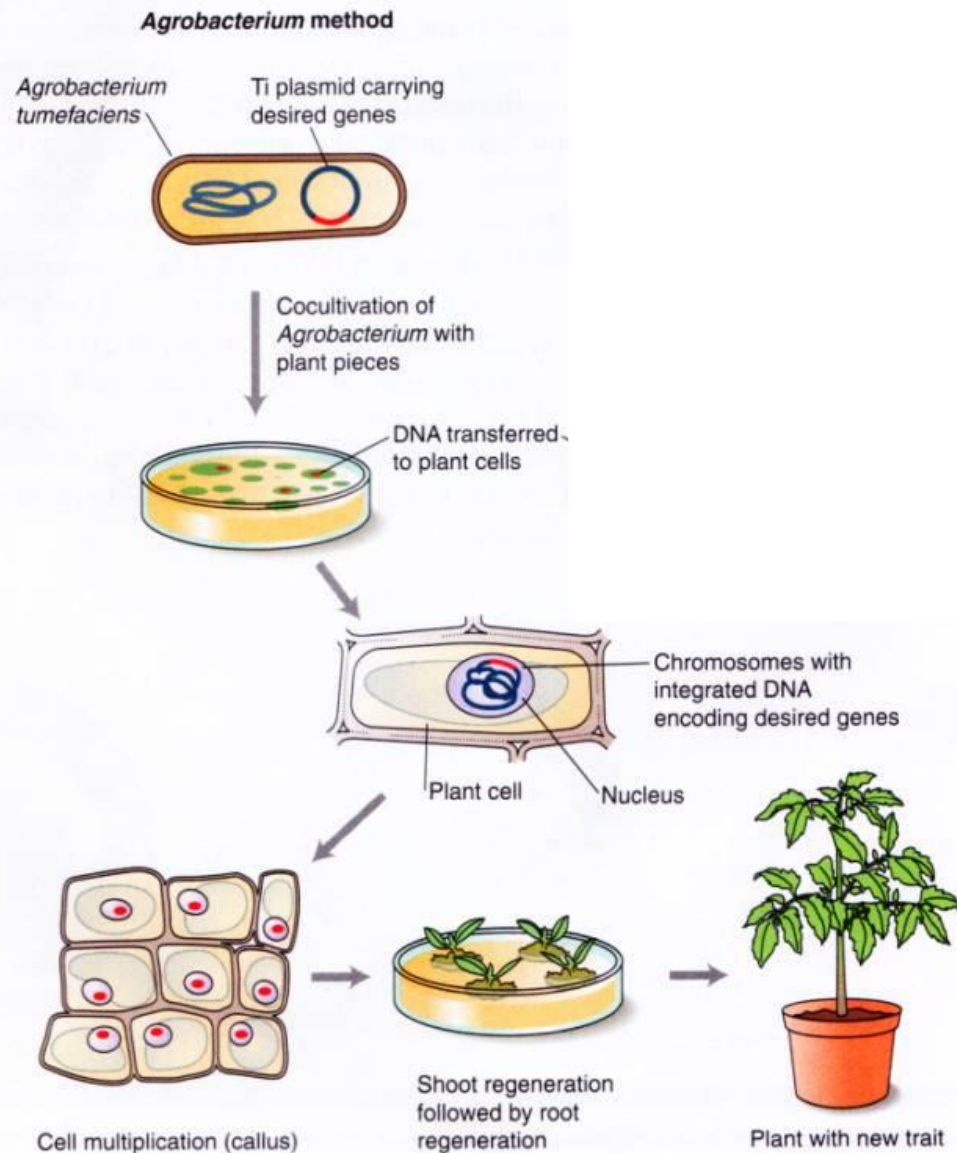


Genetic engineering through agrobacterium tumefaciens



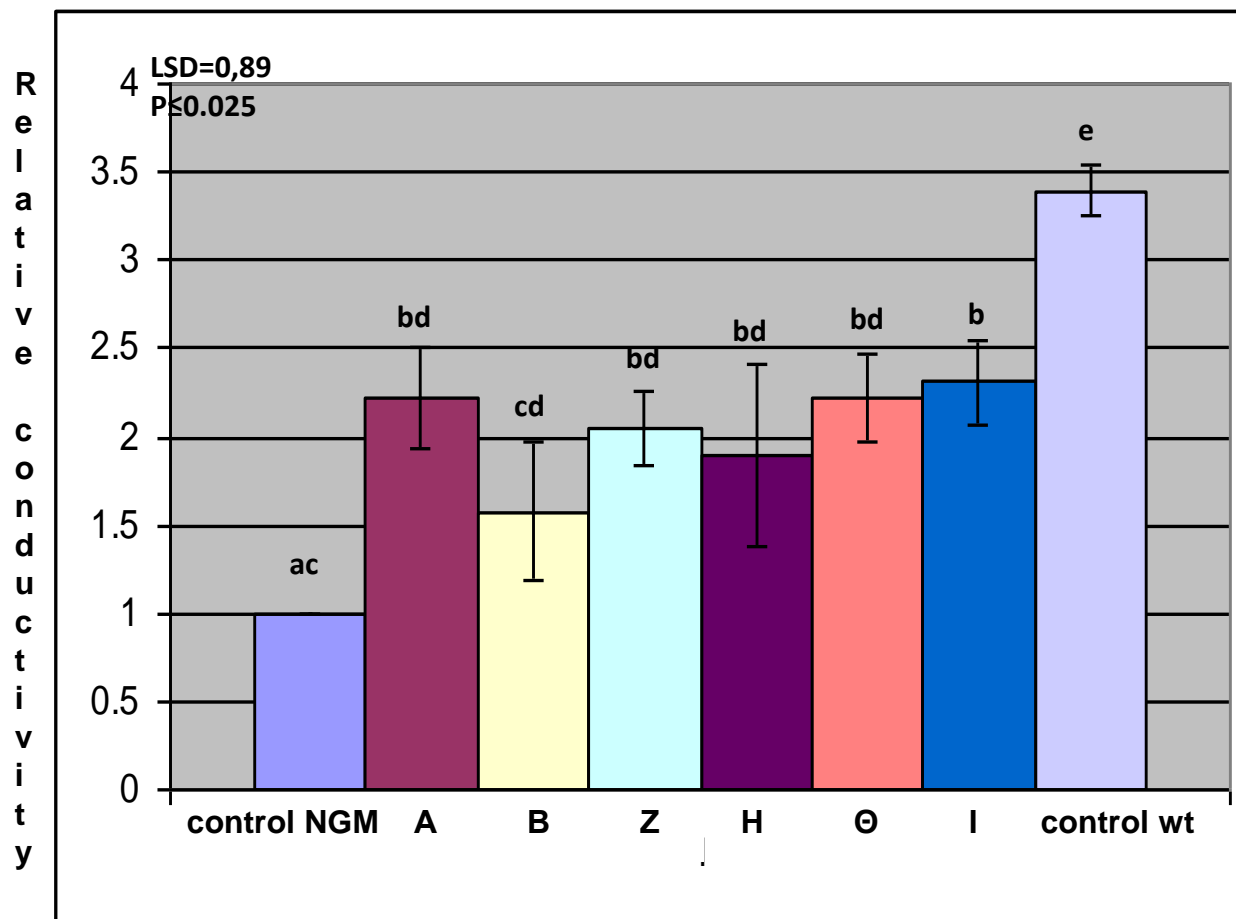


Γενετική τροποποίηση φυτών





Relative resistance of transgenic plants



Available online at www.sciencedirect.com

ScienceDirect

Biomolecular Engineering 24 (2007) 245–251

**Biomolecular
Engineering**

www.elsevier.com/locate/genengbi

Expression of the yeast *cpd1* gene in tobacco confers resistance
to the fungal toxin cercosporin

Madesis Panagiotis^a, Kalantidis Kritonas^a, Nianiou Obeidat Irini^a, Chatzidimitriou Kiriaki^a,
Panopoulos Nicolaos^a, Tsaftaris Athanasios^{a,c,*}





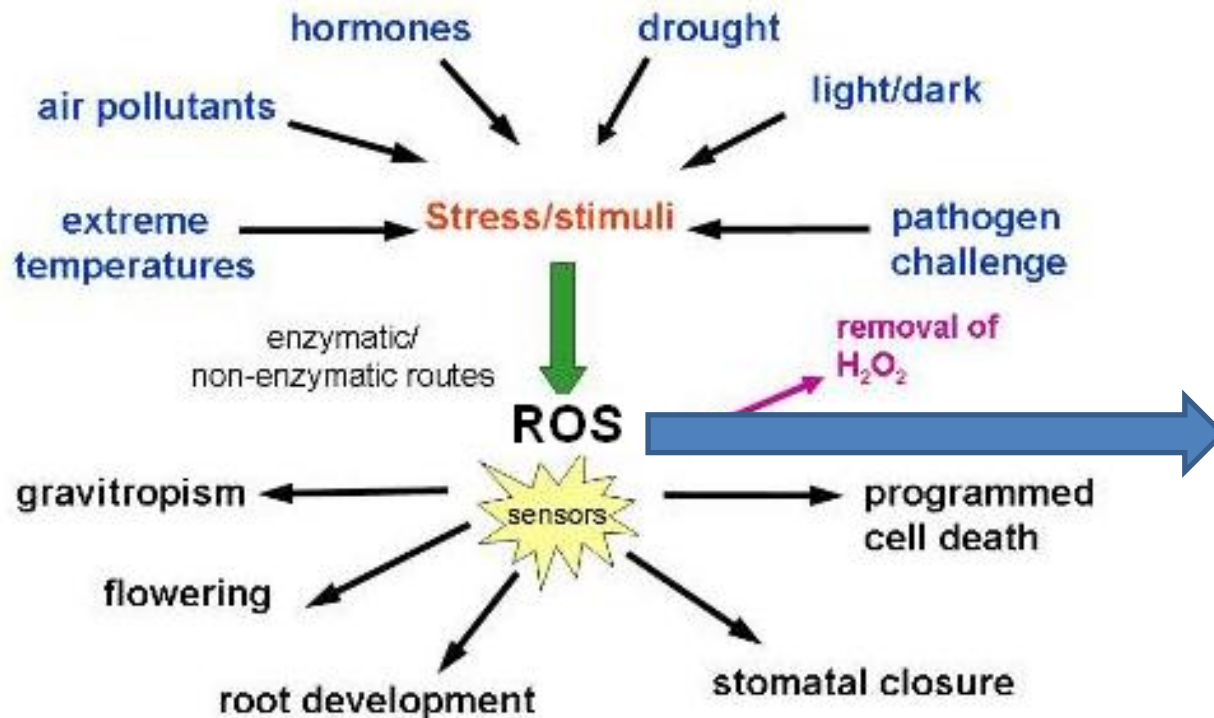
Developing stress tolerant plants





Oxidative stress

ROS: (O_2^{*-} , H_2O_2 , OH^{*-} , 1O_2)





Oxidative stress

Addressing ROS



Natural Defense :

- Avoid
- Plant Morphology

Chemical Defense:

- Vitamins
- Flavonoids
- carotenoids,
- **GSH**

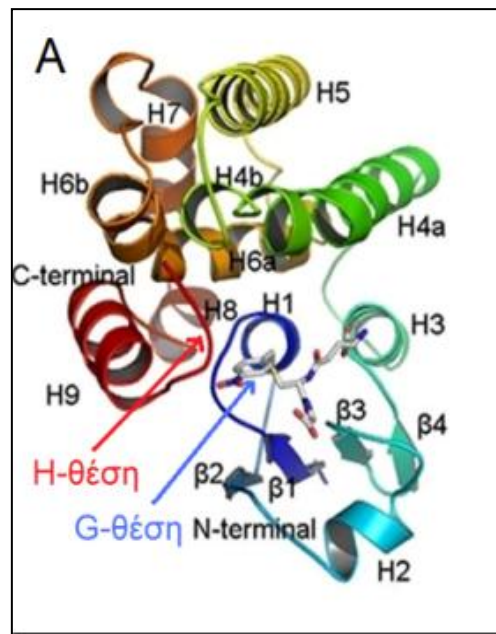
Enzyme Defense:

- Superoxide dismutase SOD,
- Catalase CAT,
- ascorbate peroxidase APX
- glutathione reductase, GR
- dehydroascorbate reductase, DHAR
- υπεροξειδάσες της γλουταθειόνης (GPX)
- **glutathione transferase, GST**





Structure of GST



Current Chemical Biology, 2011, 5, 000-000

1

Structure and Antioxidant Catalytic Function of Plant Glutathione Transferases

Evangelia Chronopoulou¹, Irene Axarli¹, Irini Nianiou-Obeidat², Panagiotis Madesis³, Athanasios Tsaftaris^{2,3} and Nikolaos E. Labrou^{1*}



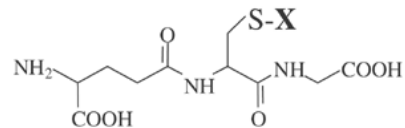


Function of GST



Glutathione

GST



Glutathione-S-Conjugate

Current Chemical Biology, 2011, 5, 000-000

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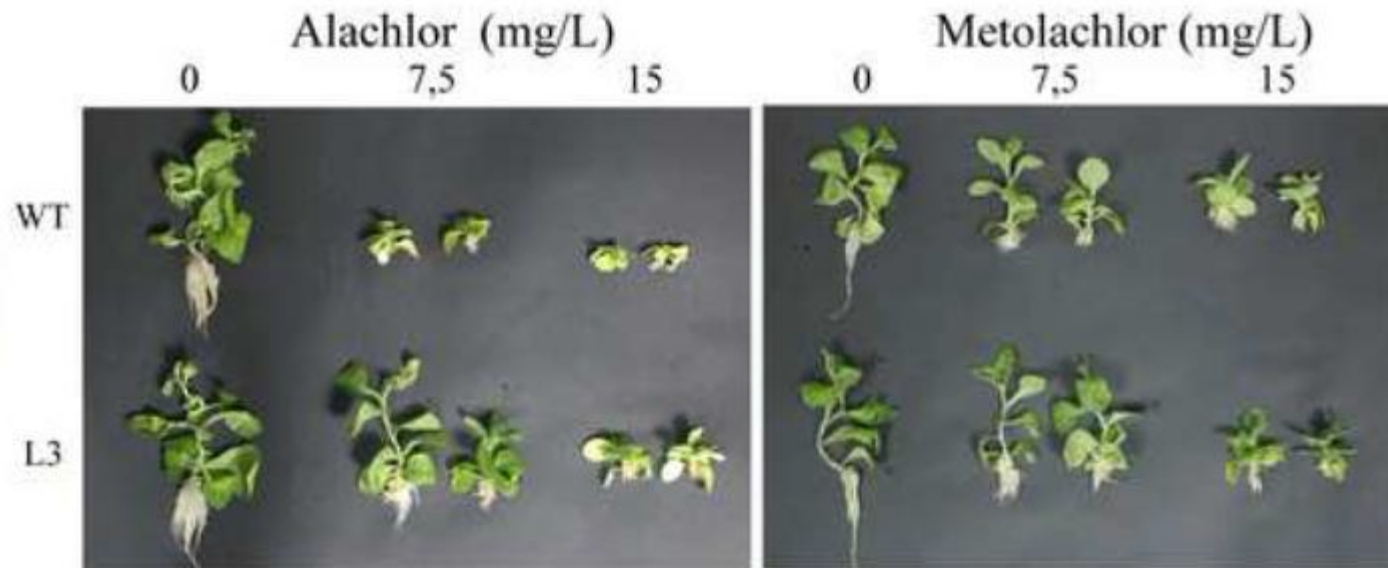
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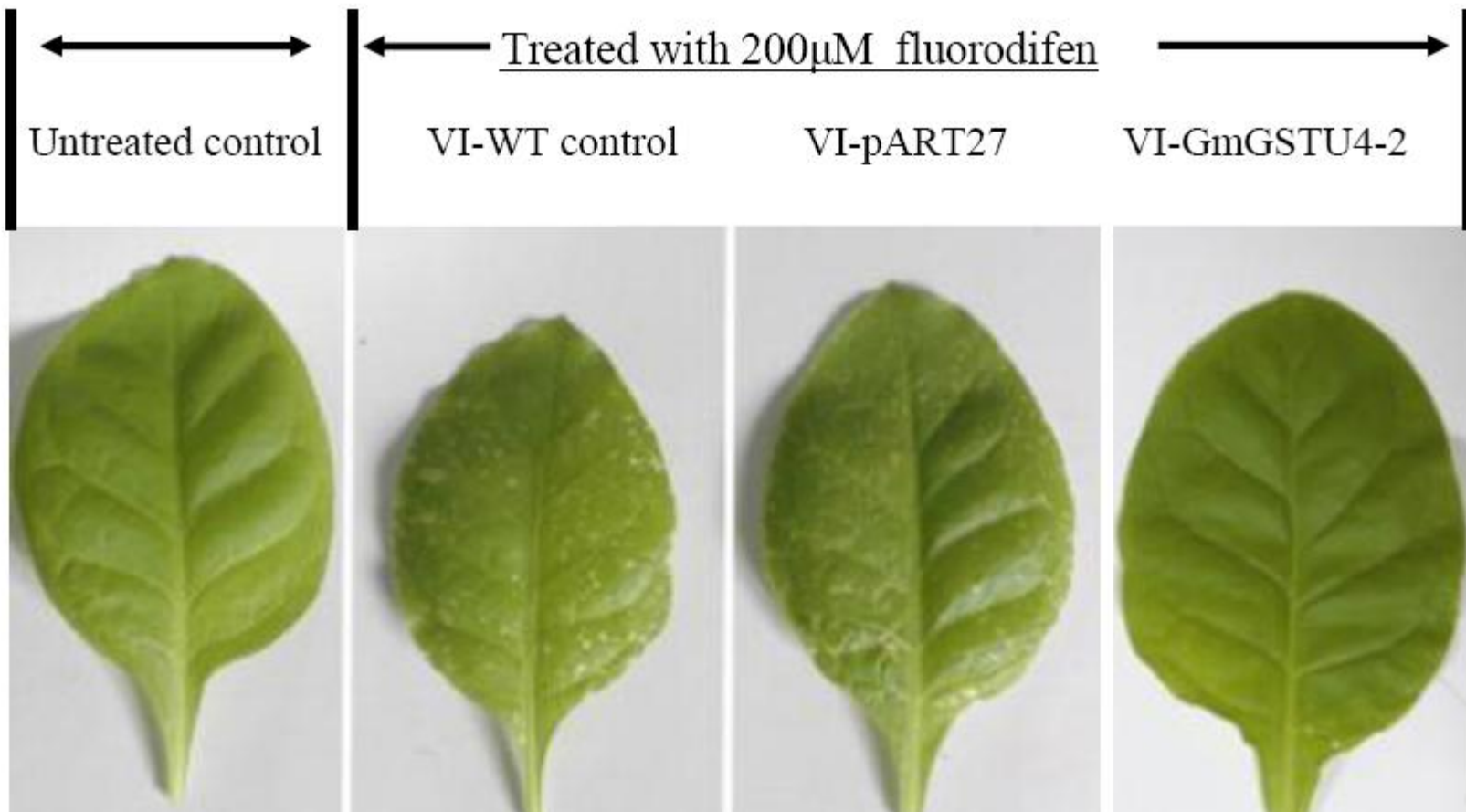


Alachlor resistance through *gst4* over-expression in tobacco



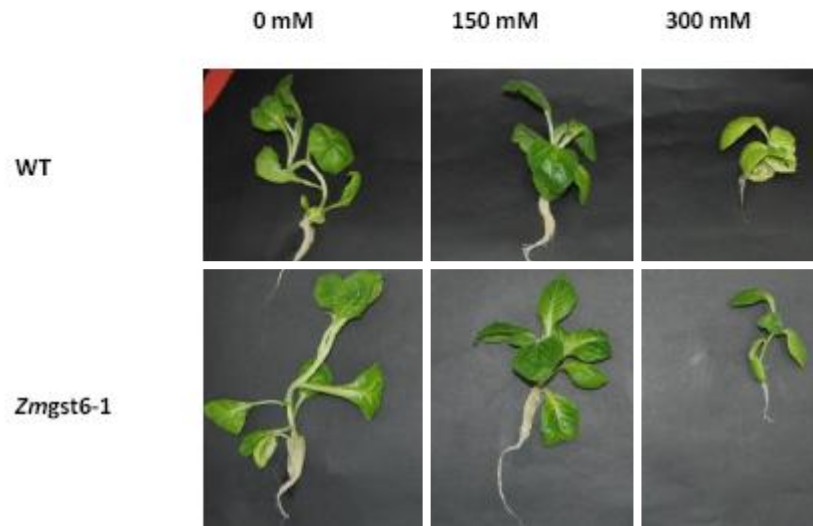


Fluorodifen resistance through *gst4* over-expression in tobacco





Salt resistance



NaCl



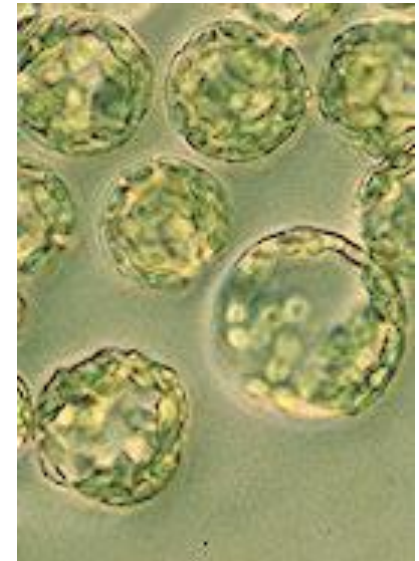
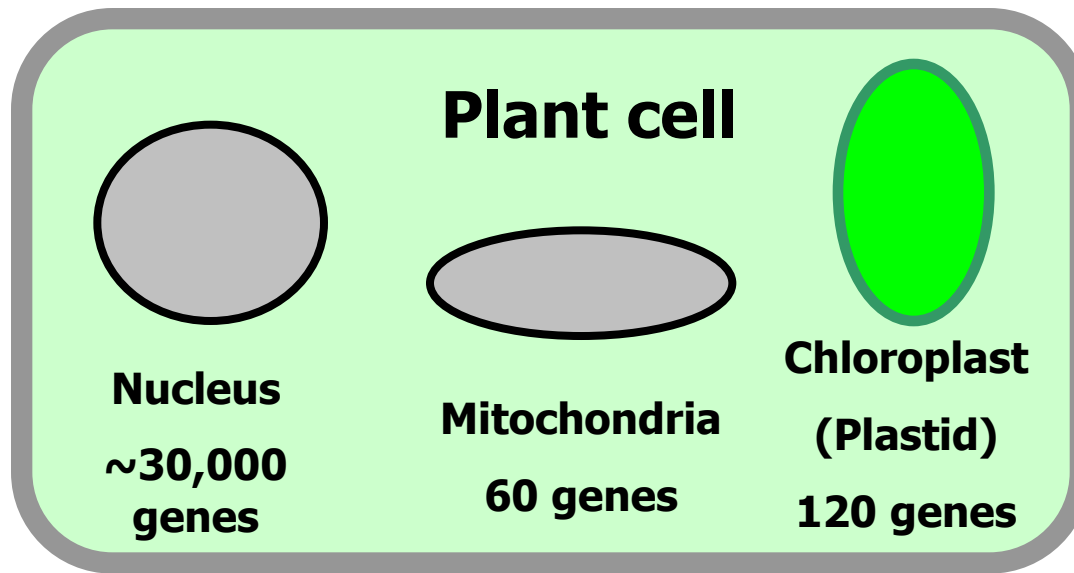


Chloroplast genetic engineering





Plants contain multiple genomes

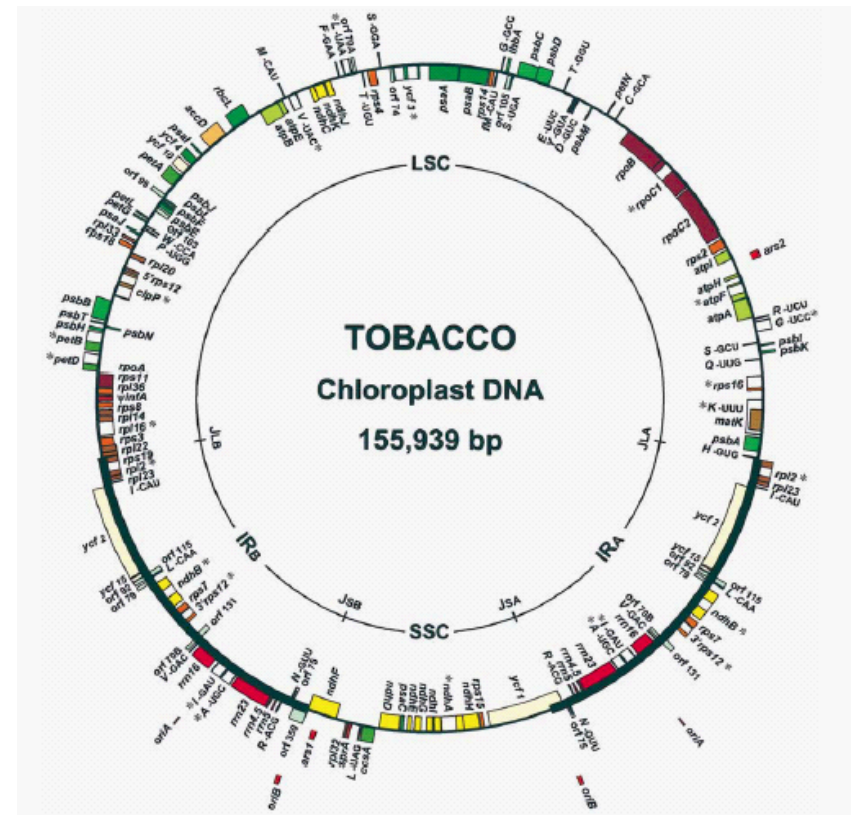
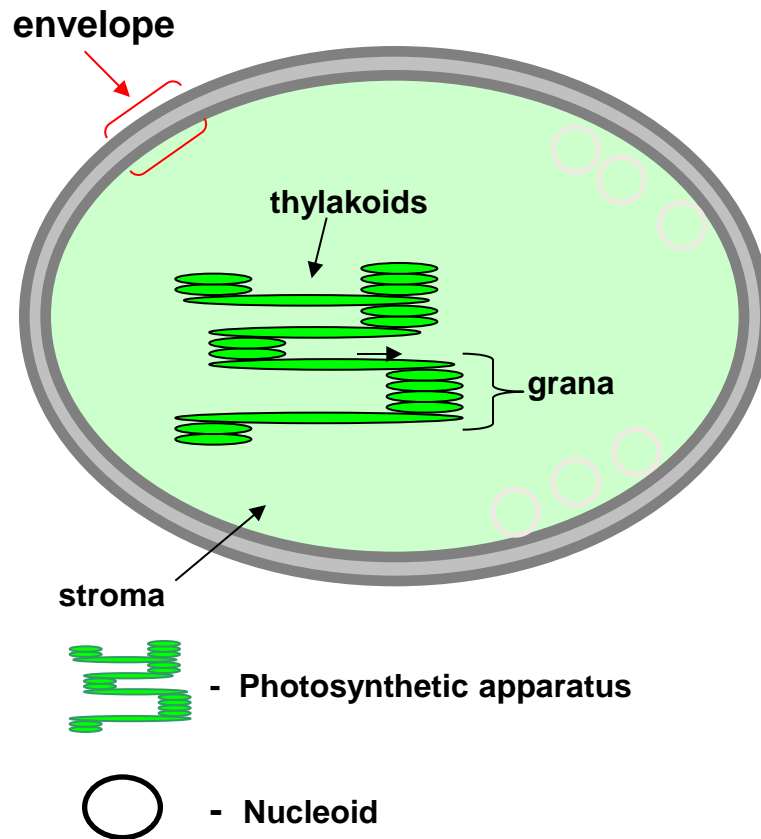


- Three genomes in all plant cells
- 100 chloroplasts per cell
- 100 genomes per chloroplast (10,000 per cell)



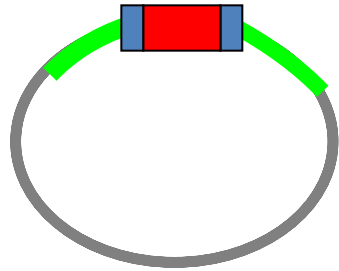


Chloroplast structure

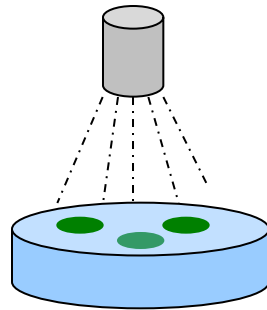




Tobacco chloroplast transformation



Plasmid stock purified and
used for particle bombardment
of leaf tissue



Three cycles of
regeneration from
leaf tissue produces
homoplasmic plants

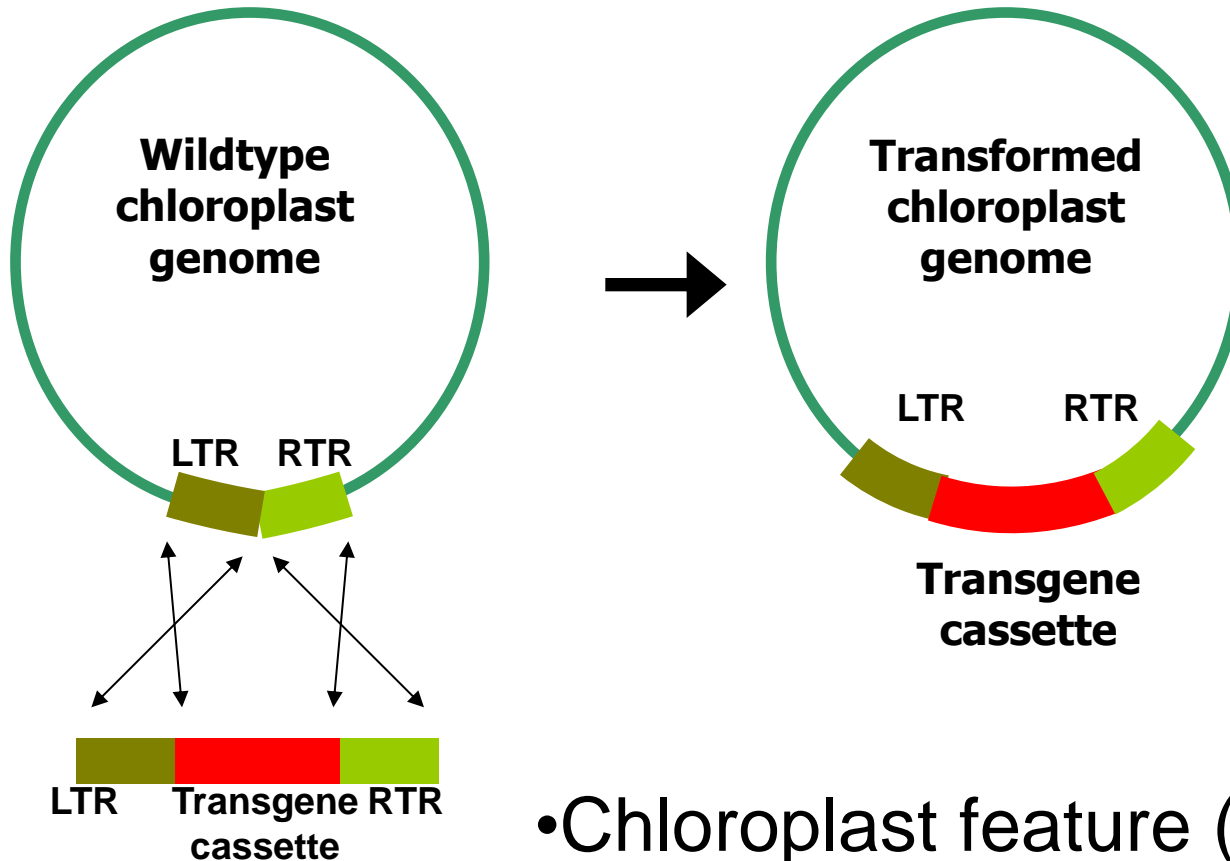


Plants regenerated from
leaf tissue under antibiotic
selection





Transgene insertion proceeds by homologous recombination

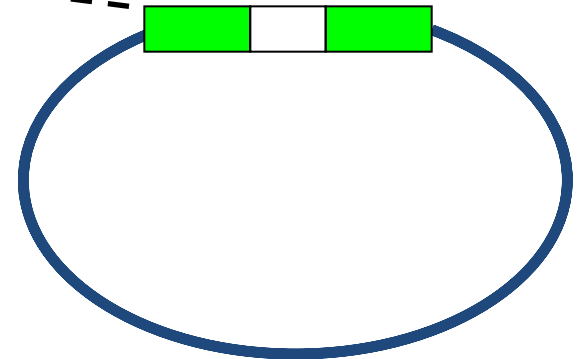
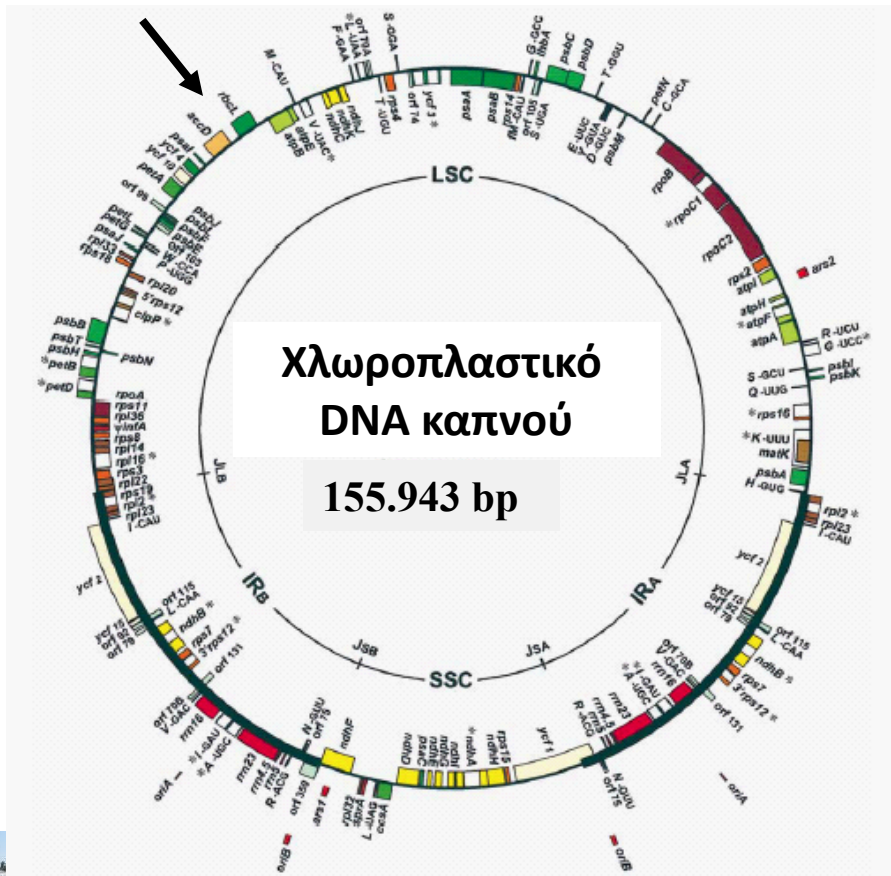


- Chloroplast feature (not nuclear)
- Targeted and stable integration





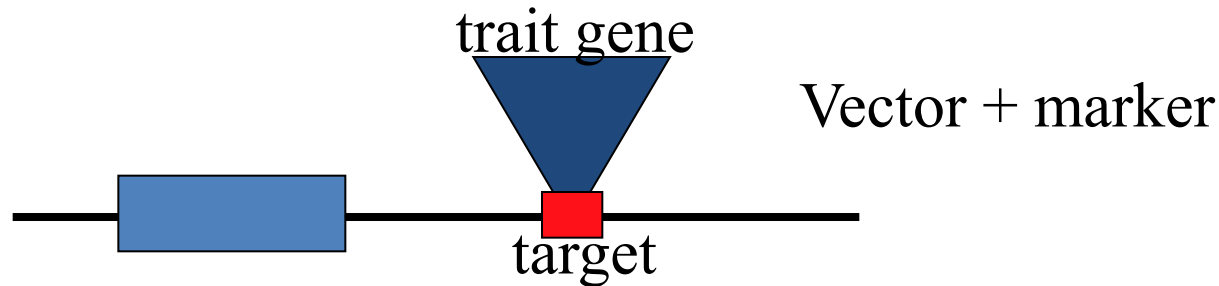
Vector structure





Clean gene transformation

- Precise integration without vector sequences or plant marker genes

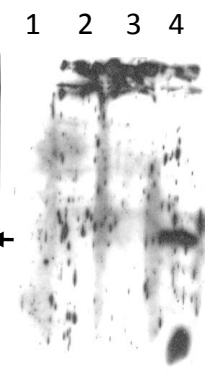
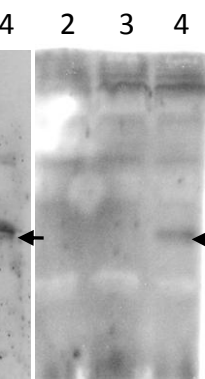
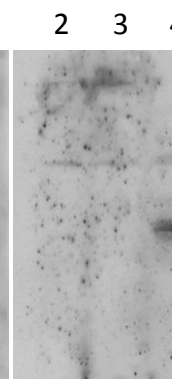
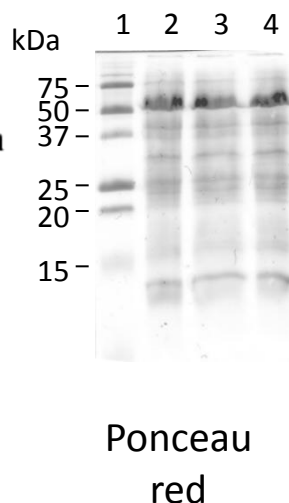
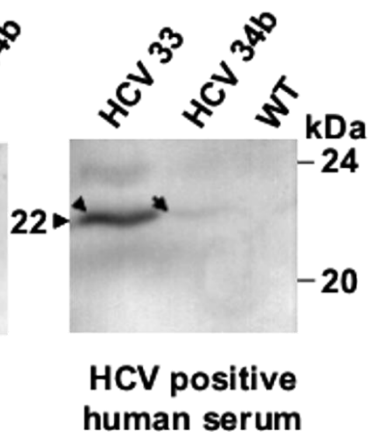
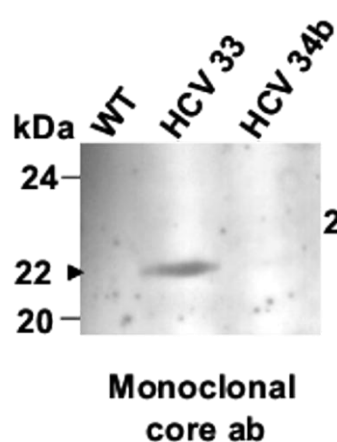
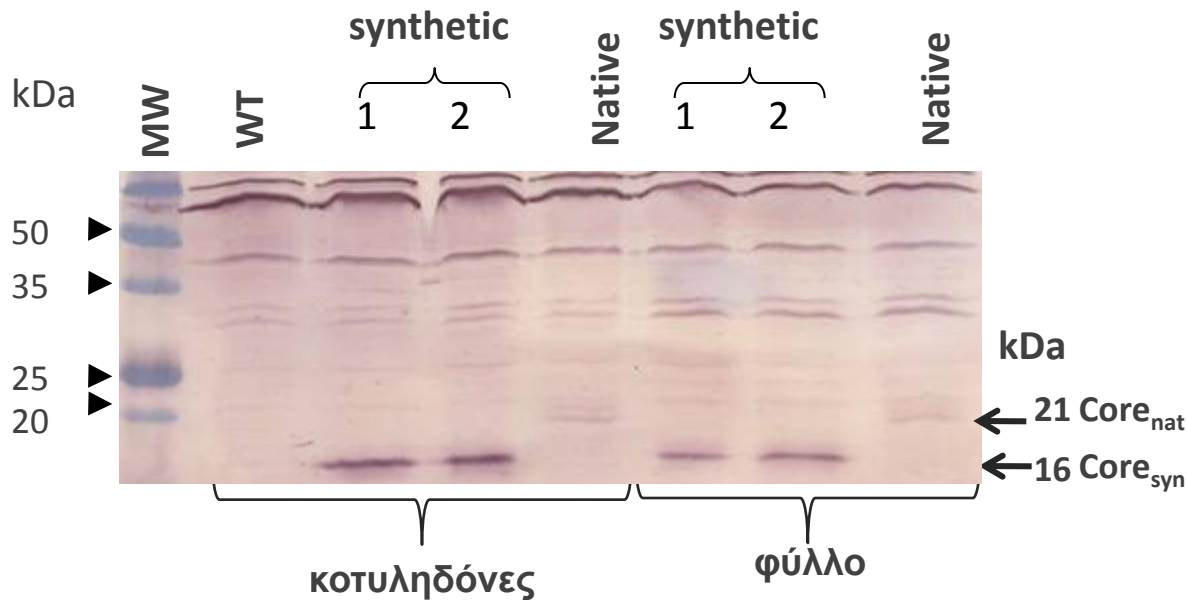


- Success of transgenic plants
 - Not only trait genes
 - improved transformation technologies





HCV core protein over-expression



Lanes

1. MW
2. WT
3. core^{nat}
4. core^{syn}

kDa

16





Thank you
for your attention

