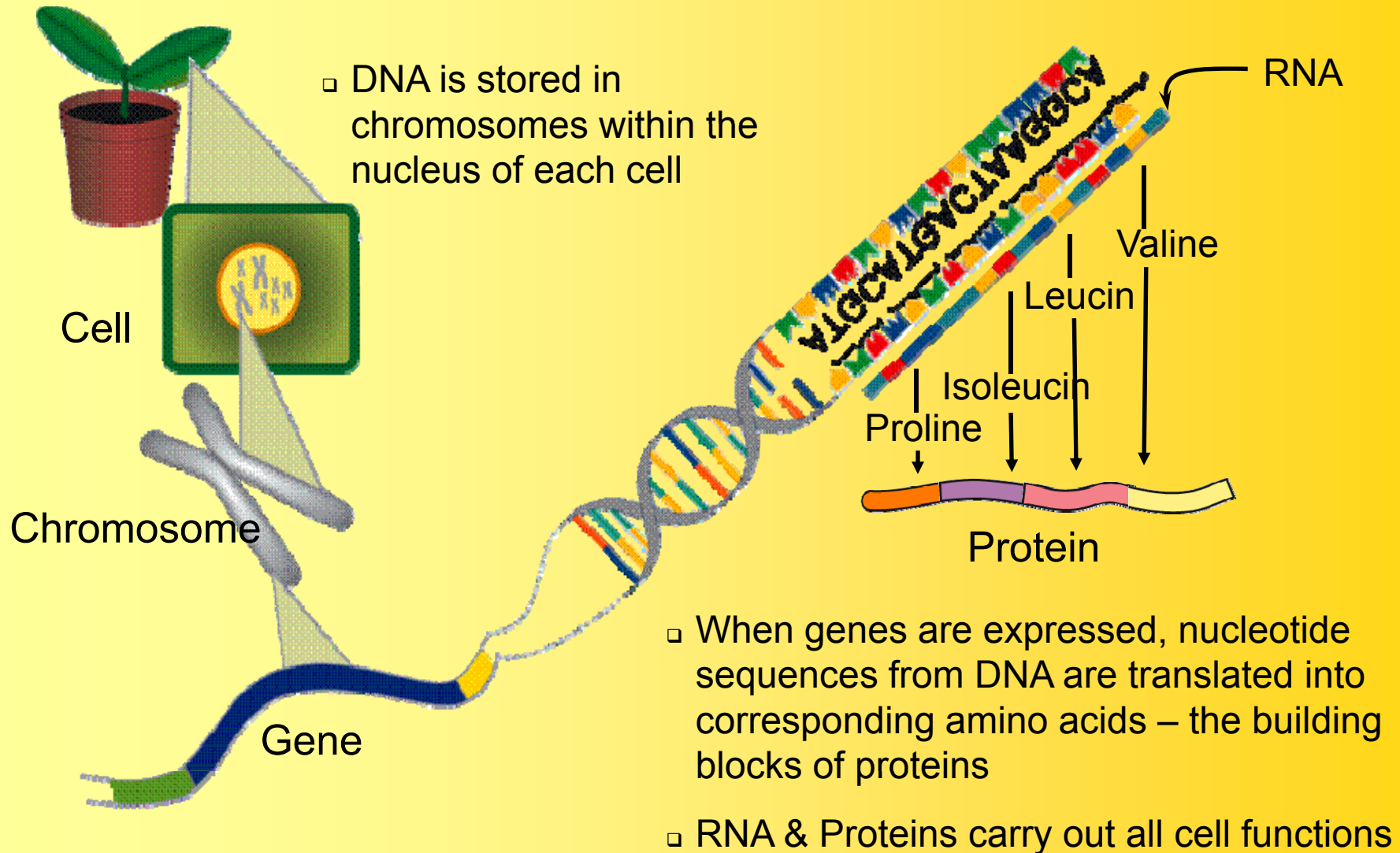


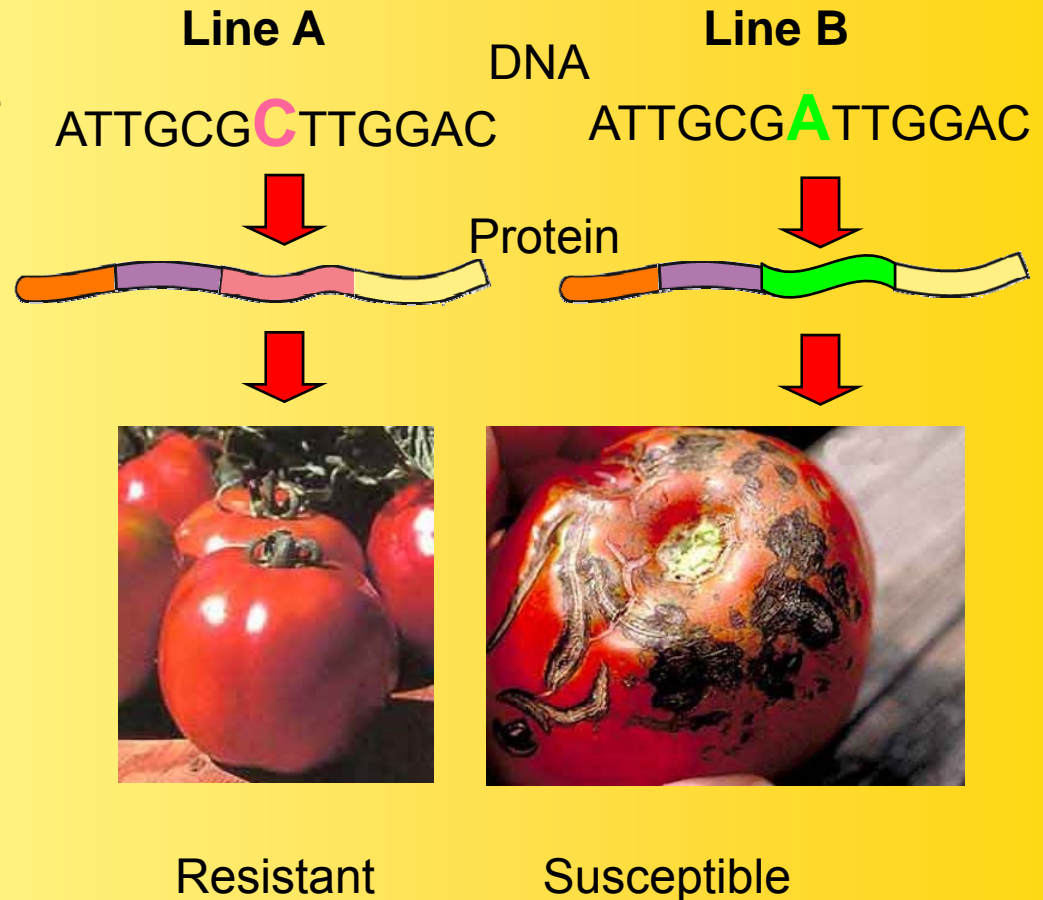
A brief introduction to Marker-Assisted Breeding





How Genotype Effects Phenotype

- Mutations can occur in DNA (e.g. cytosine to adenine or C to A)
- Some mutations cause a change in the amino acid coding at the protein level
- The new protein may have an altered function and the plant a different phenotype



Genetic markers are specific DNA sequence differences that can be identified through biochemical assays. There are two main types:

Simple Sequence Repeats (SSR)

- Usually consist of di or tri nucleotide repeats
- Also known as microsatellites
- Variation occurs in the number of repeats
- The difference in the length of each fragment can be measured
- Co-dominant marker system (more informative when dealing with heterozygotes)

Line A GTGA GT GT TGGC

Line B GTGA GT GT GT TGGC

Line C GTGA GT GT GT GT TGGC

Single Nucleotide Polymorphism (SNP)

- Based on the allelic variation of single nucleotides
- Co-dominant marker system
- Most prolific of all marker types
- Very efficient and inexpensive to use once developed

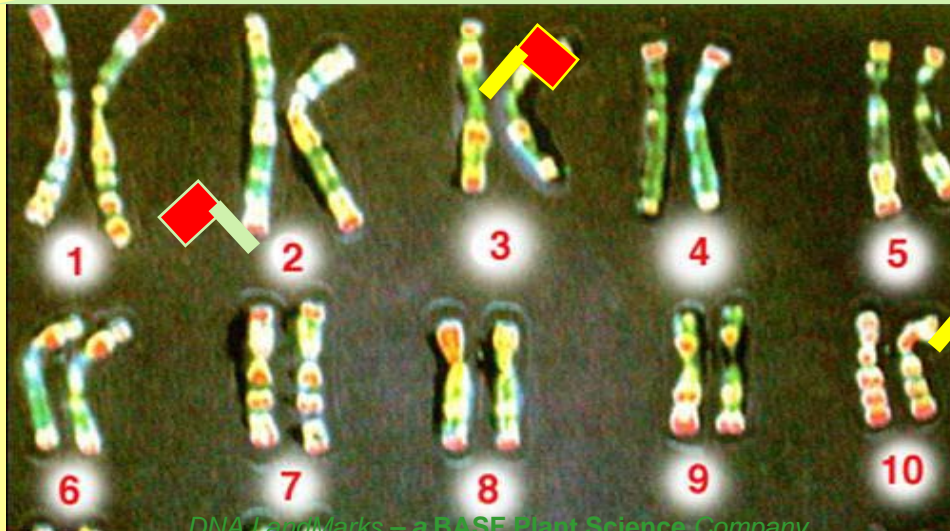
Line A CCTGTAA TGGTACATT

Line B CCTGTAA CGGTACATT

How Genetic Markers Track Genotype



- DNA markers can occur anywhere across the genome



DNA LandMarks – a BASF Plant Science Company

Chromosome picture

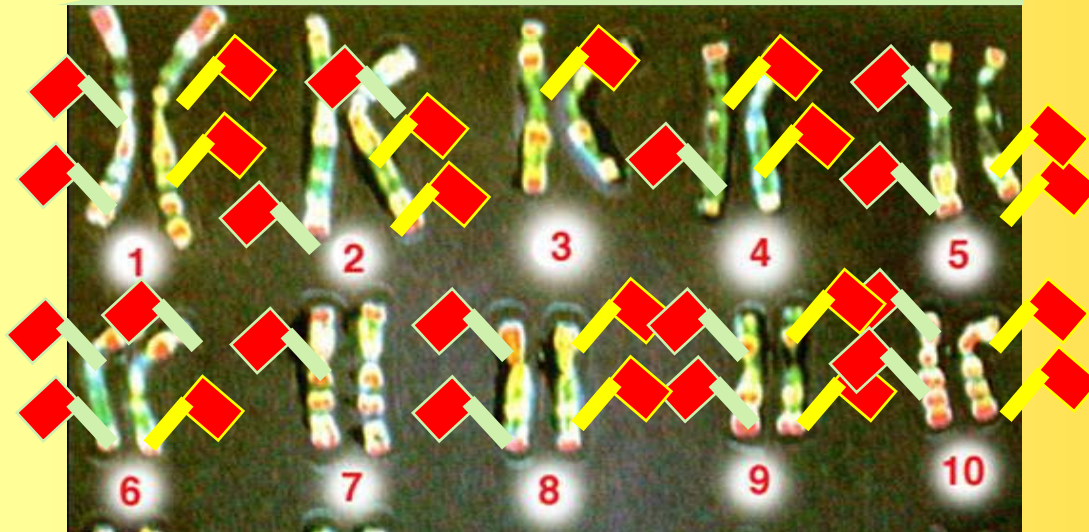
Raven, PH and GB Johnson 1991.

Understanding Biology. 2nd Edition. Mosby.
St. Louis.

How Genetic Markers Track Genotype



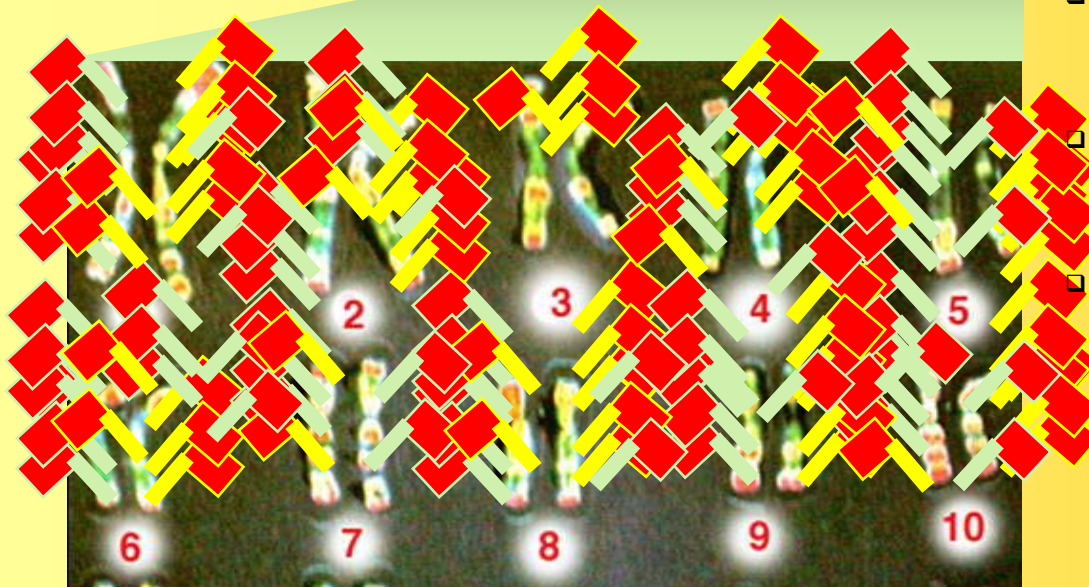
- DNA markers can occur anywhere across the genome
- The more markers you have, the more useful they become



Chromosome picture

Raven, PH and GB Johnson 1991.
Understanding Biology. 2nd Edition. Mosby.
St. Louis.

How Genetic Markers Track Genotype



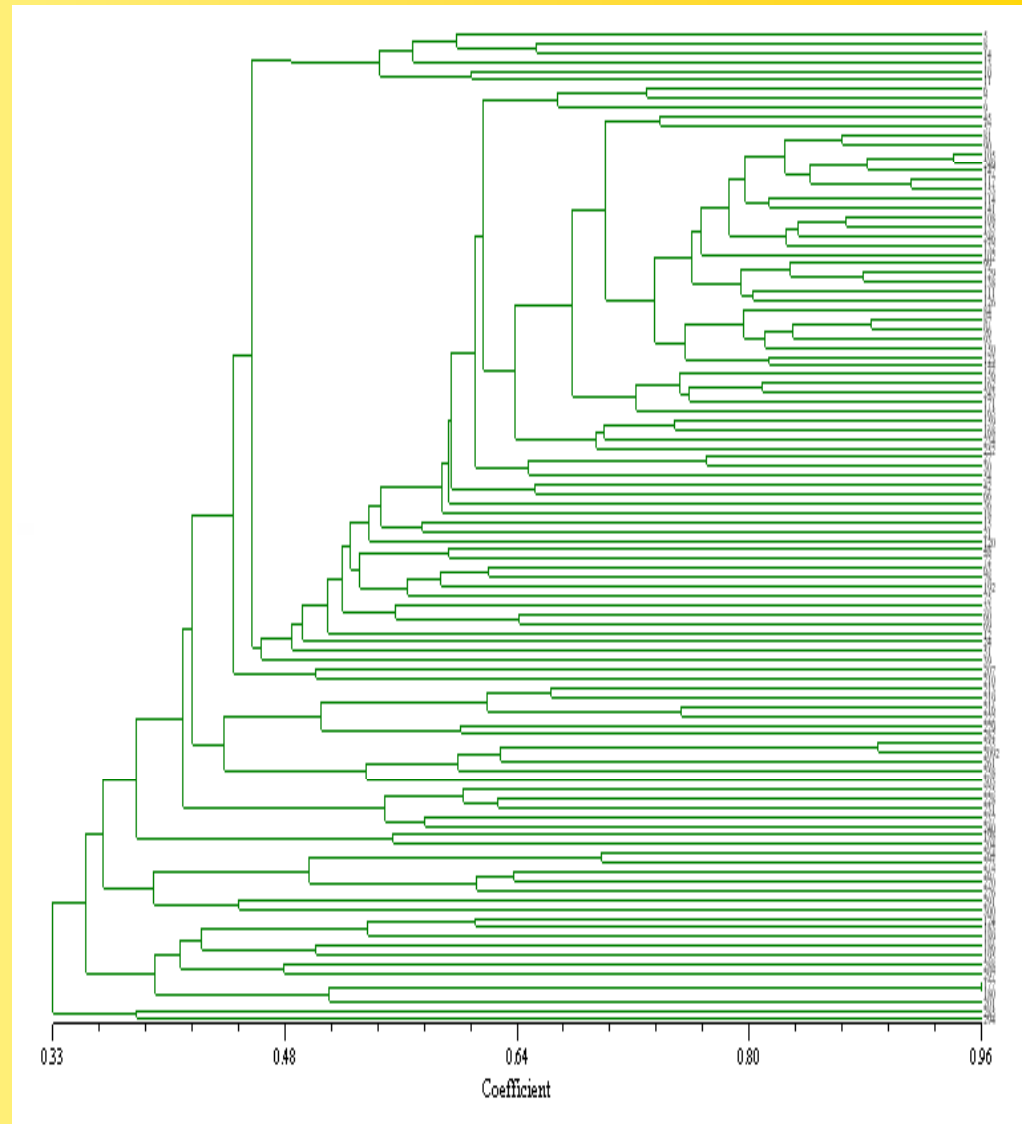
- DNA markers can occur anywhere across the genome
- The more markers you have, the more useful they become
- The ultimate goal is a “saturated” genetic map
- This allows you to find markers closely linked to any gene of interest

Chromosome picture

Raven, PH and GB Johnson 1991.
Understanding Biology. 2nd Edition. Mosby.
St. Louis.

DNA Fingerprinting

- Use a standard set of genetic markers evenly distributed across the genome
- Fingerprint each line with the same standard set of markers
- Creates a genetic “scorecard” that can be used to plan crosses
- Also widely used for quality control and protection against variety infringement



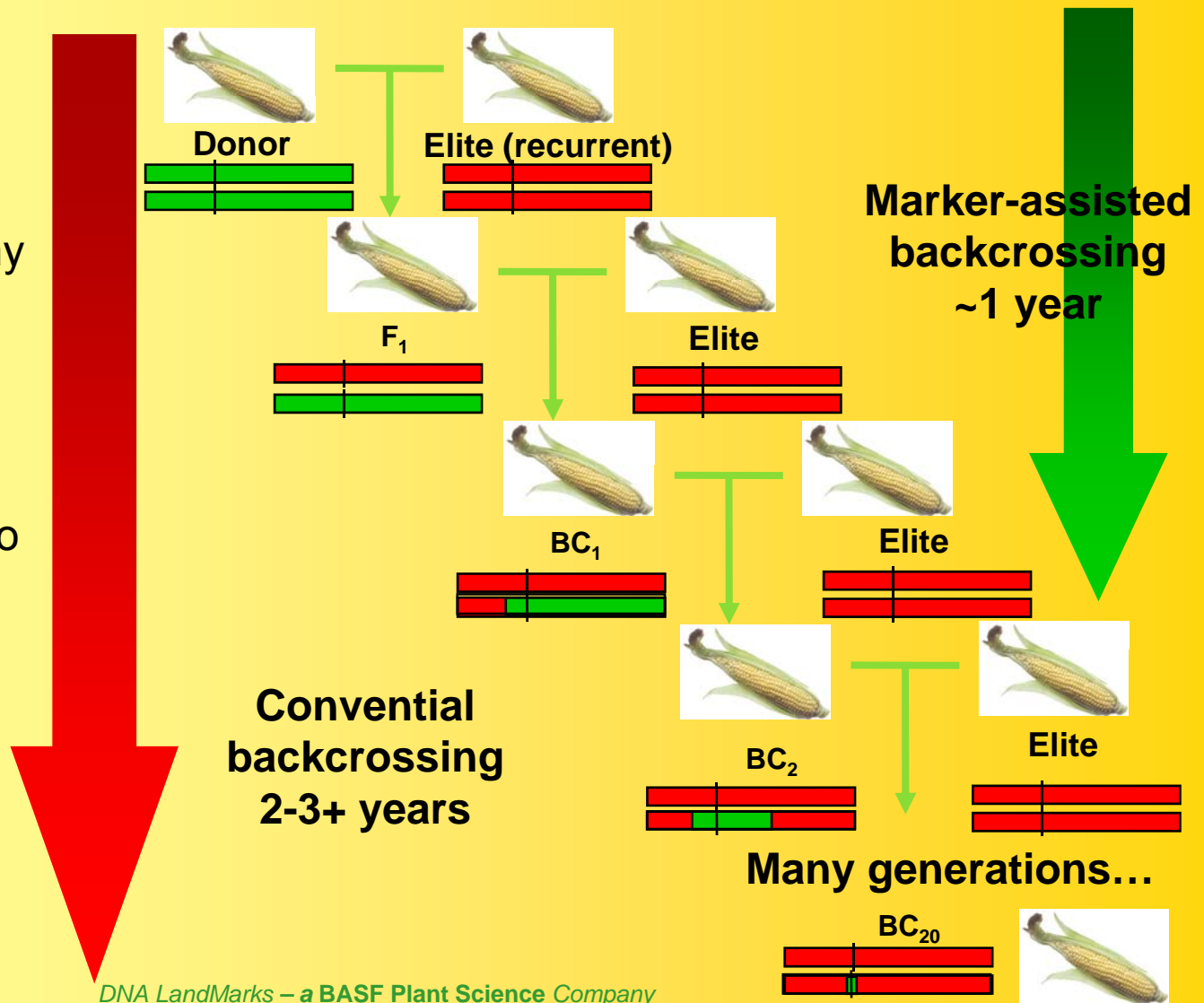
Marker-assisted selection

- Track traits of interest over generations using markers that are tightly linked to the trait's gene
- Most widely used application of genetic markers in plant breeding
- Possible to screen for single traits, multigenic traits (QTL) and to pyramid multiple desirable traits into a single variety
- Eliminates environmental effects
- Accelerates selection process
- Allows selection of traits that are difficult to evaluate phenotypically



Marker-Assisted Backcrossing

- Uses markers to compare BC progeny to the recurrent parent (RC)
- Analysis identifies rare progeny with very high similarity to RC
- Possible to find BC₂ progeny with ~95% similarity to the RC



Conclusion

Markers have become an essential and affordable tool for a competitive breeding program:

- Examine your germplasm resources at the genomic level
- Determine the components of multigenic traits through QTL mapping
- Manage complex agronomic traits
- Pyramid multiple traits into elite germplasm through selection and accelerated backcrossing
- Ensure quality control at the genetic level
- Protect your genetic investment once it is in the marketplace

Get the best varieties to market faster!