



SCST Genetics Super Workshop

SNP Testing



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What is a SNP?

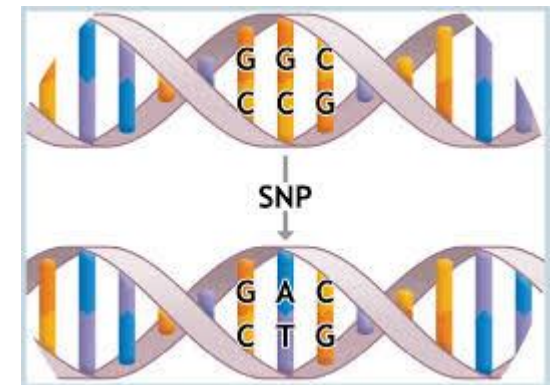
Single Nucleotide Polymorphism (SNP)

// SNP is a substitution of a single nucleotide at a specific position on the genome.

Variety 1: ATGCCGTAGTACATTTGTATGCTAATAGCG

Variety 2: ATGCCGTAGTACAGTGTATGCTAATAGCG

- // Needs to be contained in at least 1% of the species population
- // Usually, biallelic
- // Can occur in non-coding or coding regions
- // Caused by mutation and natural selection
- // Are inherited
- // 100Ks – 1Ms of SNPs can exist in whole genomes
- // May or may not be trait-associated



<https://www.socmucimm.org/resources/news-media/single-nucleotide-polymorphism-snp-allele-frequency-dna-pools/>



Identifying SNPs

Genome Sequencing is used to find SNP locations

// Populations

// Breeding varieties

Reference -ATGCCGTAGTACA[T/G]TGTATGCTAATAGCG....
Variety 1 –ATGCCGTAGTACATTGTATGCTAATAGCG....
Variety 2 –ATGCCGTAGTACAGTGTATGCTAATAGCG....
Variety 3 –ATGCCGTAGTACATTGTATGCTAATAGCG....
Variety 4 –ATGCCGTAGTACAGTGTATGCTAATAGCG....
Variety 5 –ATGCCGTAGTACAGTGTATGCTAATAGCG....

// Genome-Wide Association Studies (GWAS)

// Aim to identify associations between SNPs and phenotypic traits

// Public Databases

// Research institutions



Creating a SNP Panel

Informativity vs. Cost

Determine the Purpose: Line identification vs Line confirmation

- // Breeding Programs
 - // Early in the process
 - // Ensure advancing correct lines
 - // Small number of samples
 - // More markers = high genome representation
- // Quality Control Programs
 - // Later in the process
 - // Quality check to confirm production processes
 - // Larger number of samples
 - // Less markers = lower cost

Things to Consider

- // Informativity
 - // The ability to differentiate between lines within a population
- // Allele distribution by marker
 - // PIC-value
- // Number of markers needed
- // Genome representation
- // Inbred vs Hybrid informativity



PIC Value

Polymorphism Information Content (PIC)

- // Allele distribution by marker
- // Ideally, 50:50 distribution of alleles
 - // PIC value: ~0.5 = high diversity
 - // PIC value: > 0.25 and <0.5 = intermediate diversity
 - // PIC value: <0.25 = low diversity
- // $PIC = 1 - (allele\ 1^2 + allele\ 2^2)$

	Mk1	Mk2	Mk3
Allele 1	50	90	10
Allele 2	50	10	90
Total lines	100	100	100
Allele 1 + Allele 2	100	100	100
Allele 1%	0.50	0.90	0.10
Allele 2%	0.50	0.10	0.90
Allele 1% ²	0.25	0.81	0.01
Allele 2% ²	0.25	0.01	0.81
PIC : 1-(Allele 1² + Allele 2²)	0.50	0.18	0.18



SNP Marker Panel Informativity

- // Informativity
 - // The ability to differentiate between lines within a population
- // Example:
 - // 10 varieties but only 5 unique genotypes
 - // With this panel if varieties 1 and 3 were accidentally mixed together or swapped, the test wouldn't be able to tell them a part - maybe we care, maybe we don't
- // Understanding testing scope will help determine correct panel informativity

	Mk1	Mk2	Mk3	Mk4	Mk5
Var1	Allele 1	Allele 2	Allele 1	Allele 2	Allele 1
Var2	Allele 1	Allele 2	Allele 1	Allele 2	Allele 2
Var3	Allele 1	Allele 2	Allele 1	Allele 2	Allele 1
Var4	Allele 1	Allele 2	Allele 1	Allele 2	Allele 2
Var5	Allele 1	Allele 2	Allele 1	Allele 2	Allele 1
Var6	Allele 1	Allele 2	Allele 2	Allele 1	Allele 2
Var7	Allele 1	Allele 2	Allele 2	Allele 1	Allele 1
Var8	Allele 1	Allele 2	Allele 2	Allele 1	Allele 2
Var9	Allele 1	Allele 2	Allele 2	Allele 1	Allele 1
Var10	Allele 2	Allele 1	Allele 2	Allele 1	Allele 2
PIC : 1-(su	0.18	0.18	0.50	0.50	0.50



Technologies for SNP Marker Detection

// TaqMan

// Qualitative end-point

// Quantitative real-time

// KASP® – Kompetitive Allele-Specific PCR – LGC/Biosearch Technologies

// rhAMP® - Integrated DNA Technologies

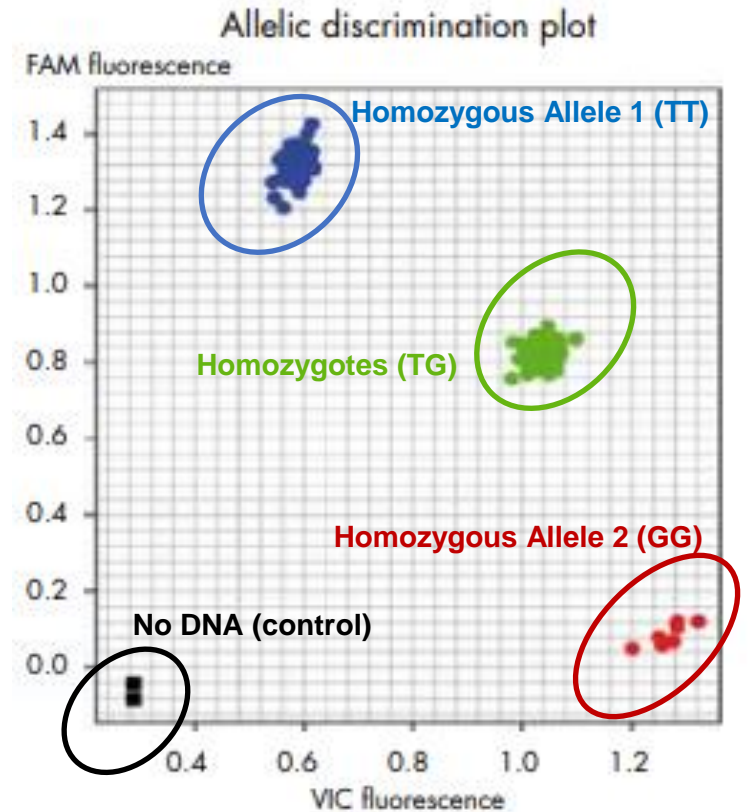
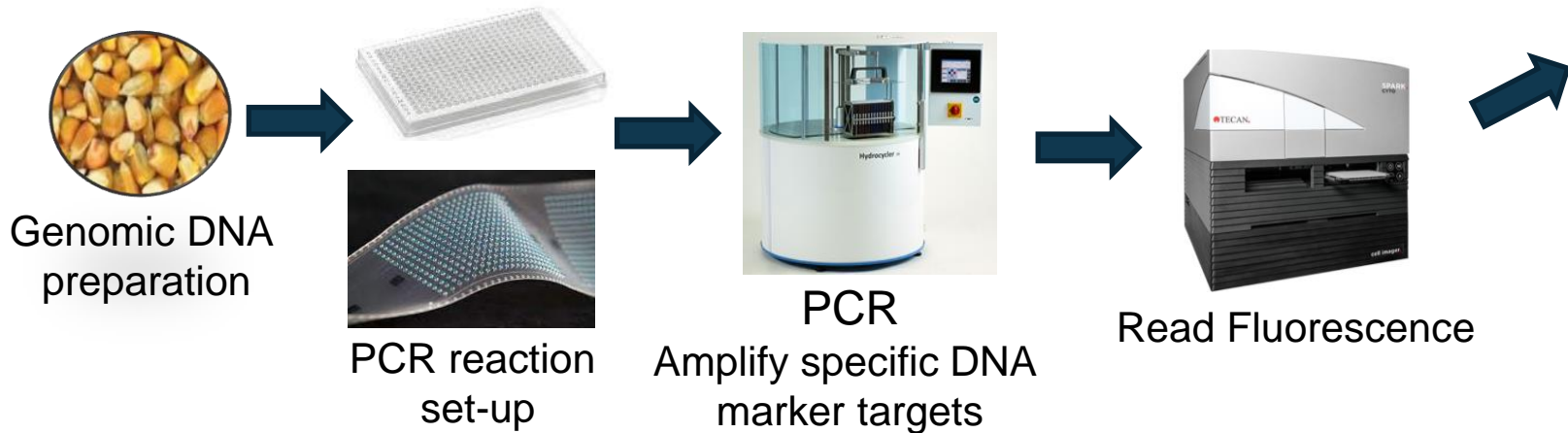
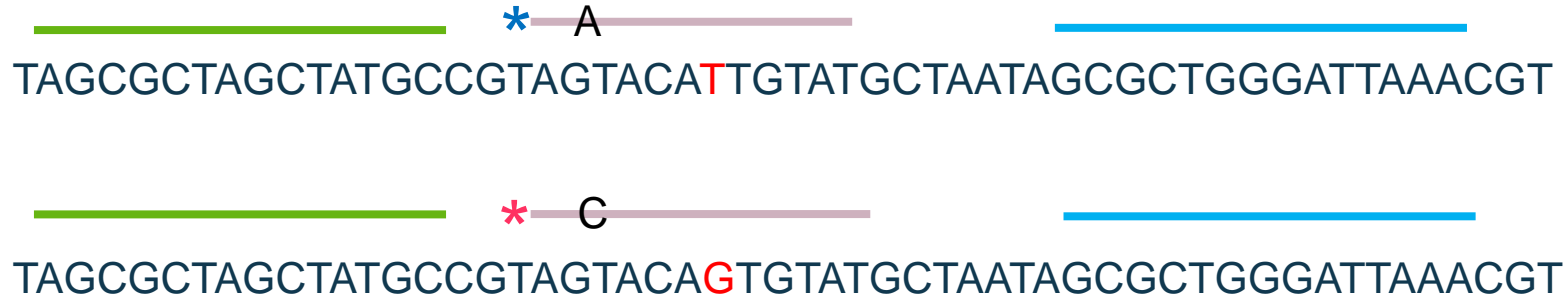
// Microarray-based Detection

// Targeted genotyping by sequence (tGBS)



Endpoint TaqMan PCR

Technologies cont.



Type-it® Fast SNP Probe PCR Kit, product insert (ver.12/2015). Qiagen

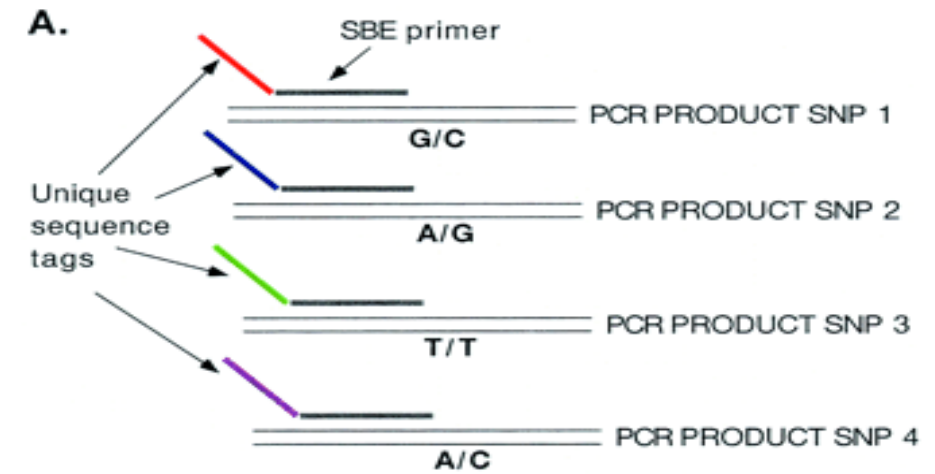


Multiplexing

Technologies cont.

- // Targeting multiple SNP locations in a single reaction or tube/well
- // Thousands of SNPs detected in a single reaction
- // Specific primers and probes for each allele

- // GeneChip – 900,000+ SNP detection



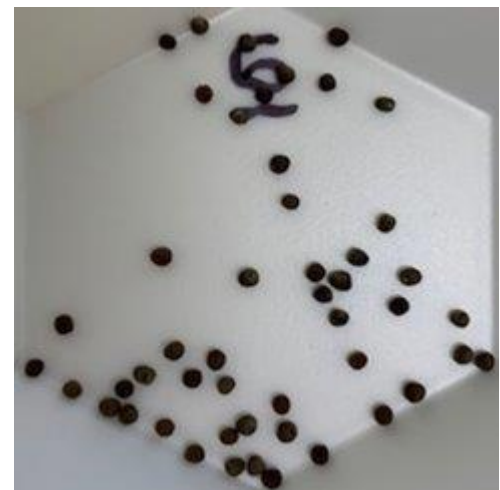
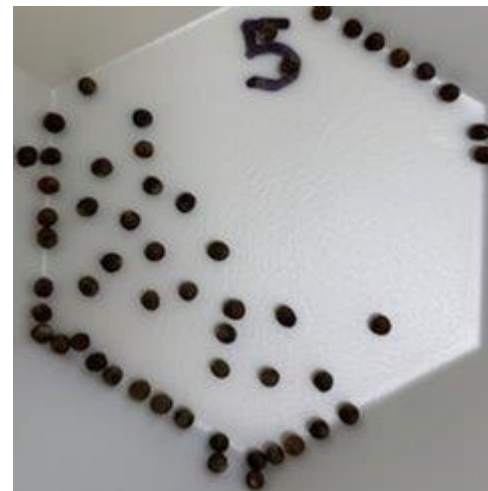
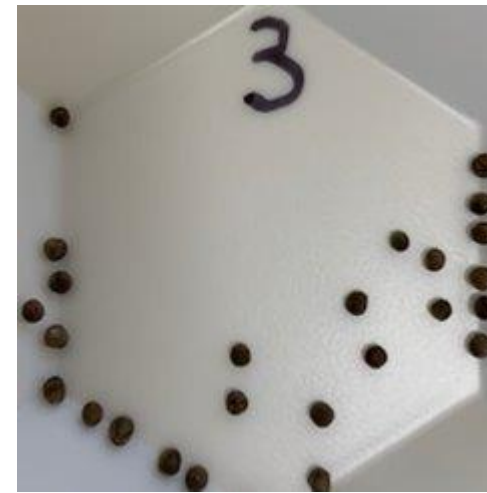
PNAS October 24, 2000 97 (22) 12164-12169; <https://doi.org/10.1073/pnas.210394597>



<https://www.bumc.bu.edu/microarray/services/affymetrix-genechip/#:~:text=Affymetrix%20GeneChip%C2%AE%20System%20for%20Genotyping&text=This%20new%20approach%20increases%20the,markers%20in%20the%20replication%20phase.>



Can You Guess the Varieties (or Crop)?



Cheddar 1



PRODUCT DETAILS		
Days to Maturity	Head Color	Head Curd
80-84 Days	Medium-orange	Dense
Head Shape	Plant Size	
Domed	Medium-large	

SNP Profile - H 1 2 H 1 1 2 H 2 1

Freedom 6



PRODUCT DETAILS		
Days to Maturity	Head Color	Head Curd
72-77 Days	Very white	Dense
Head Shape	Plant Size	
Domed	Medium	

SNP Profile - 1 1 2 1 H 1 2 1 H 1

Omphalos 4



Product Details	
Fruit Colour	Dark green
Plant Type	
Related Maturity / Picking	75 - 80 days
Fruit Size	1.5 to 1.8kgs
Fruit Shape/Skin	Round

SNP Profile - 2 H H 1 1 H 1 2 H 2

Green Voyager 2



Product Details	
Fruit Colour	Blue green
Plant Type	
Related Maturity / Picking	15 - 20 days
Fruit Size	1.4 to 1.6kgs
Fruit Shape/Skin	Round

SNP Profile - H H 1 H H H H 2 H 2

Castle Dome 3



PRODUCT DETAILS		
Relative Maturity	Bead Size	Color
Early	Fine	Green
Head Type		
Tight dome		

SNP Profile - H 2 1 2 H H 1 H H H

Ironman 5



PRODUCT DETAILS		
Relative Maturity	Season	Bead Size
Mid-late	Warm	Medium fine
Color	Head Type	Desert or Coastal
Blue green	High dome	Desert

SNP Profile - 1 H 1 H 2 1 2 1 H H



Thank you!

Questions?