



 Wisconsin Newborn Screening Laboratory

Current Testing Platforms Used to Detect One/Few Gene Mutations in Routine Newborn Screening

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Special Considerations

- Utilities
- Targeted mutation analysis
- Platforms
 - Available instrumentation
 - Workflow
 - Targeted gene sequences

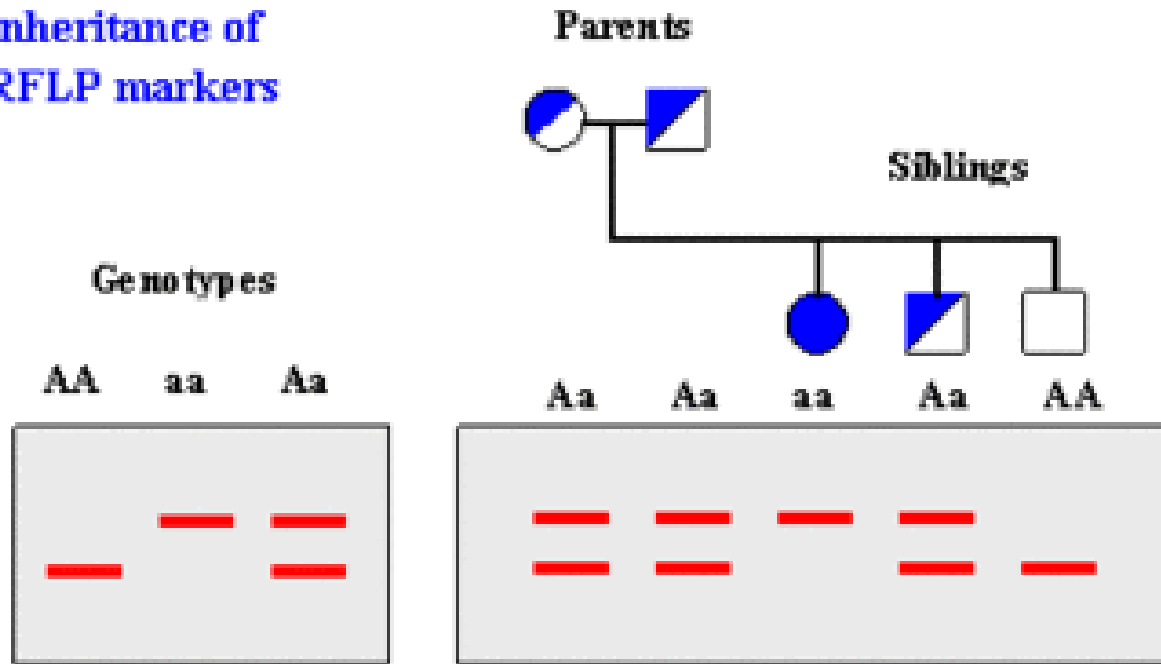
Restriction Fragment Length Polymorphism (RFLP) Analysis

<u>Enzyme</u>	<u>Recognition Site</u>
Rsa 1	.. G T A C C A T G ..
Mbo 1	.. G A T C C T A G ..
EcoR1	.. G A A T T C C T T A A G ..

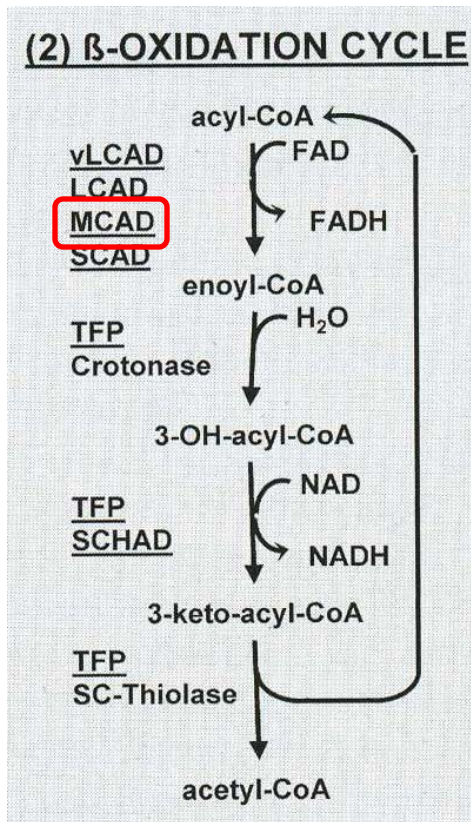
The red triangles indicate where the enzyme cuts the DNA.

Restriction Fragment Length Polymorphism (RFLP) Analysis

Inheritance of RFLP markers



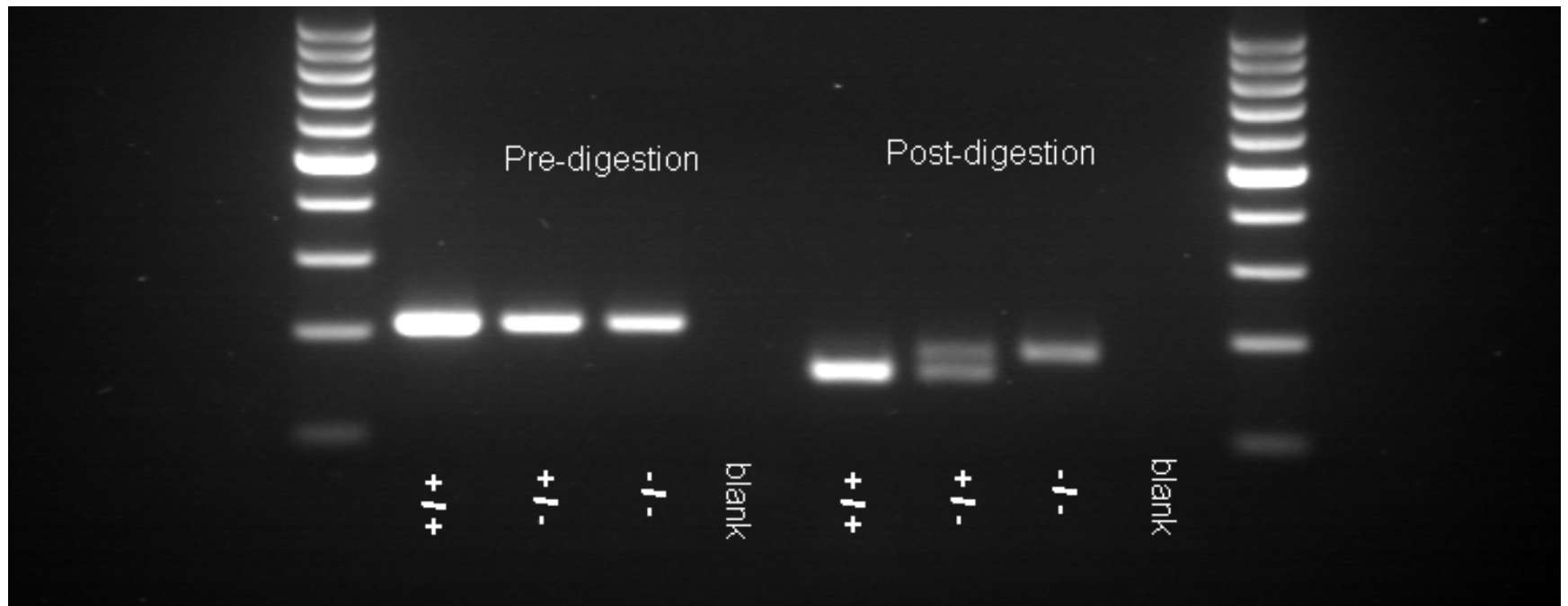
Medium-chain acyl-CoA dehydrogenase deficiency (MCADD)



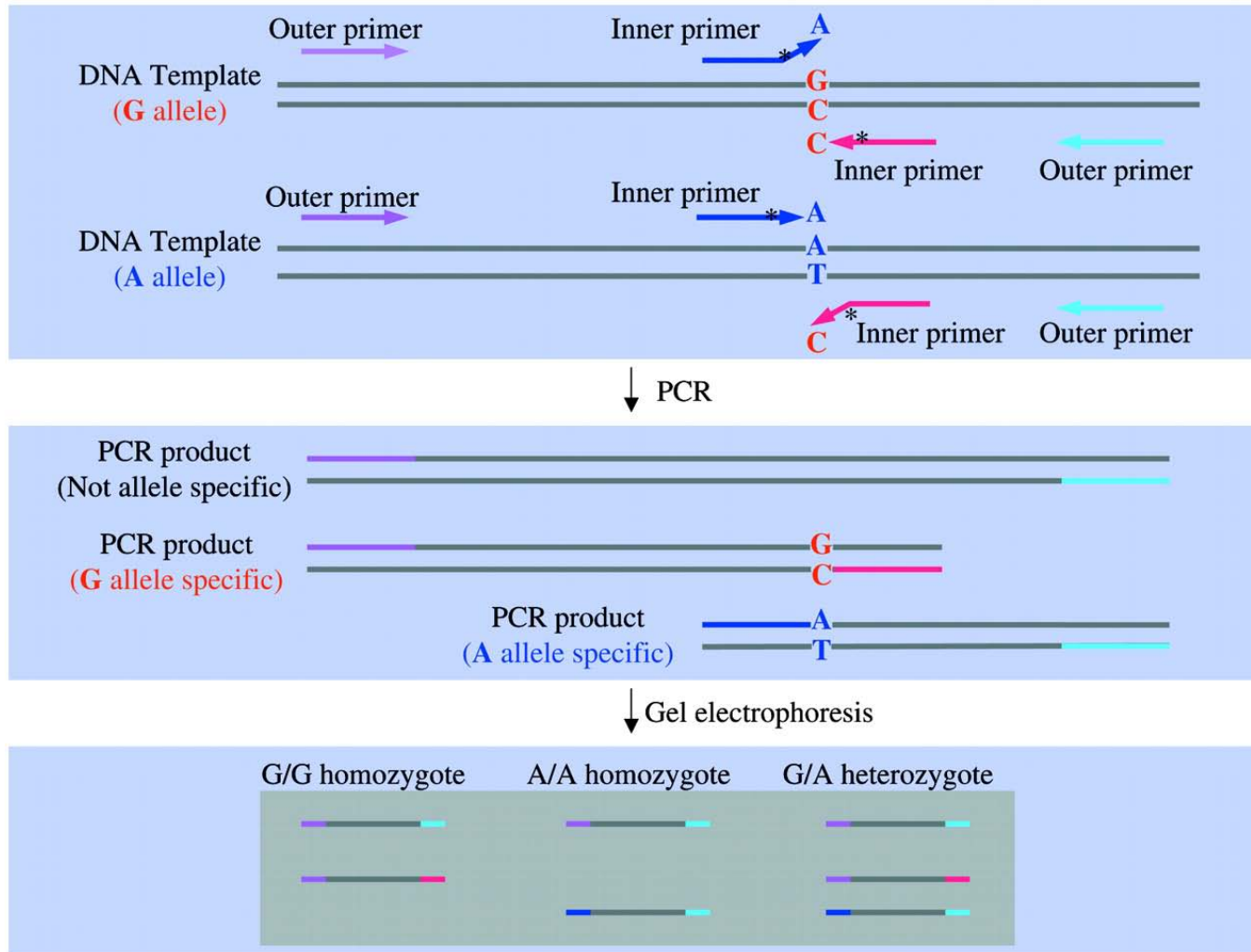
A previously healthy individual presents with:

- lethargy, seizures, and coma triggered by a common illness
- Hepatomegaly and acute liver disease
- Cardiac presentation
- Sudden and unexplained death

ACADM c.985 A>G Mutation Detected by RFLP (Nco I Digestion)



Tetra-primer ARMS-PCR



Ye, S. et al. Nucl. Acids Res. 2001 29:e88; doi:10.1093/nar/29.17.e88

Primer Design

http://cedar.genetics.soton.ac.uk/public_html/primer1.htm

Source sequence (up to 1,000 bases)

Position of SNP from start of sequence

Allele 1

Allele 2

Optimum (inner) product size

Maximum (inner) product size

Minimum (inner) product size

Maximum relative size difference of two inner products

Minimum relative size difference of two inner products

Tetra-primer ARMS-PCR Reaction

- **Reaction Mix: (25 μ l)**

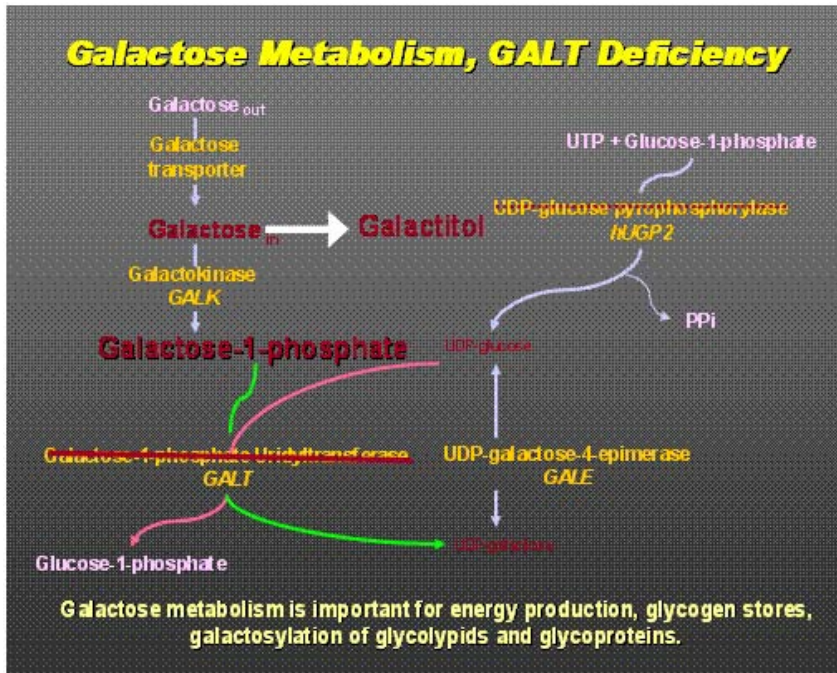
1X PCR buffer	
Forward inner primer	1.0 μ M
Reverse inner primer	1.0 μ M
Forward outer primer	0.1 μ M
Reverse outer primer	0.1 μ M
DNTPs	200 μ M
MgCl ₂	2.5 mM
<i>Taq</i> DNA polymerase	2.5 U
Genomic DNA	4 μ l

- **Thermal Cycler Condition**

1. 95°C for 5 minutes
2. 95°C for 30 second
3. 64°C for 30 second
4. 72°C for 40 second
5. repeat 2-4 for 32 cycles
6. 72°C for 2 minutes

1. The assays for different mutations are run simultaneously using the same thermal cycler conditions.
2. For N314D, inner primers concentration is 0.068 μ M, and outer primers concentration is 0.25 μ M (Rachel Lee)

Galactosemia



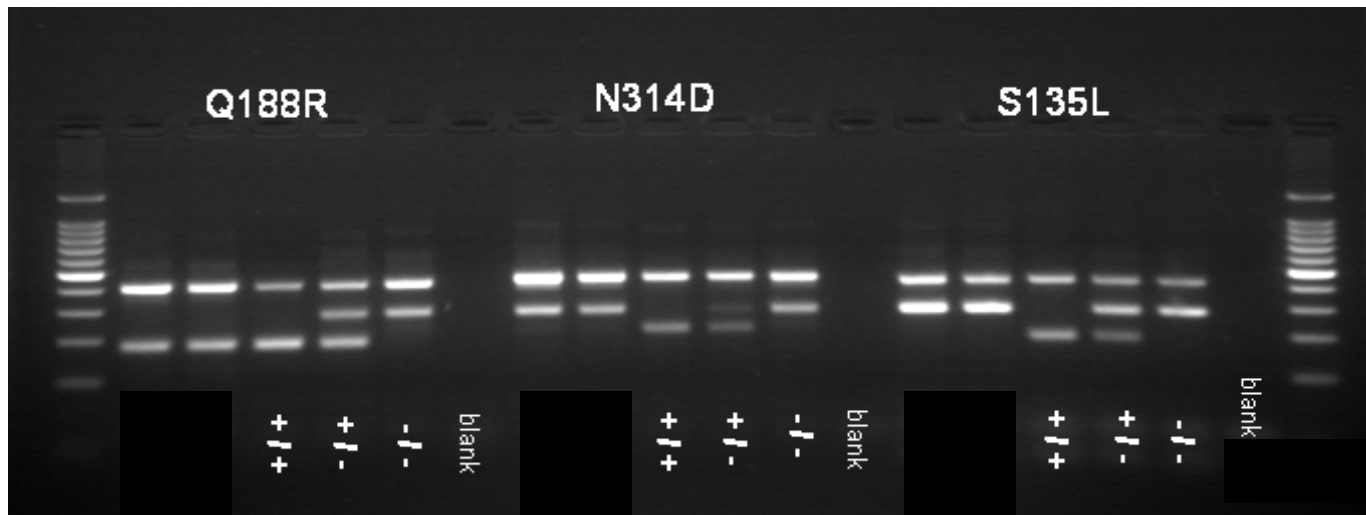
Neonates with Classic Galactosemia

Finding	Percent
Hepatocellular damage	89%
Food intolerance	76%
Failure to thrive	29%
Lethargy	16%
Seizures	1%
Sepsis	10%

Common *GALT* galactosemia (G) mutations:

[p.Gln188Arg](#), [p.Ser135Leu](#), [p.Lys285Asn](#),
[p.Leu195Pro](#), [p.Tyr209Cys](#), [p.Phe171Ser](#), [5kdel](#),
[c.253-2A>G](#)

GALT mutations Detection Using Tetra-primer ARMS-PCR



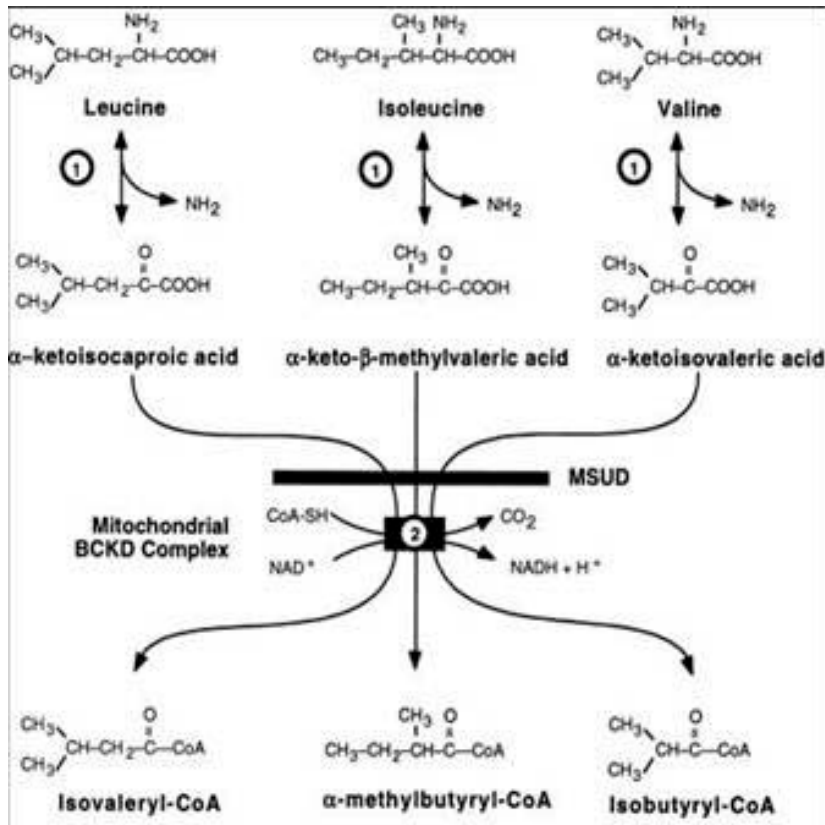
Greg Kopish

Notes:

1. Q188R—one of common galactosemia disease causing mutations.
2. N314D--Duarte (D_2) variant, and reducing enzyme activity by 25%
3. S135L is associated with a mild phenotype.
4. Newborns who are G/D heterozygotes may have a positive newborn screen

Maple Syrup Urine Disease (MSUD)

BAA Metabolism Pathway



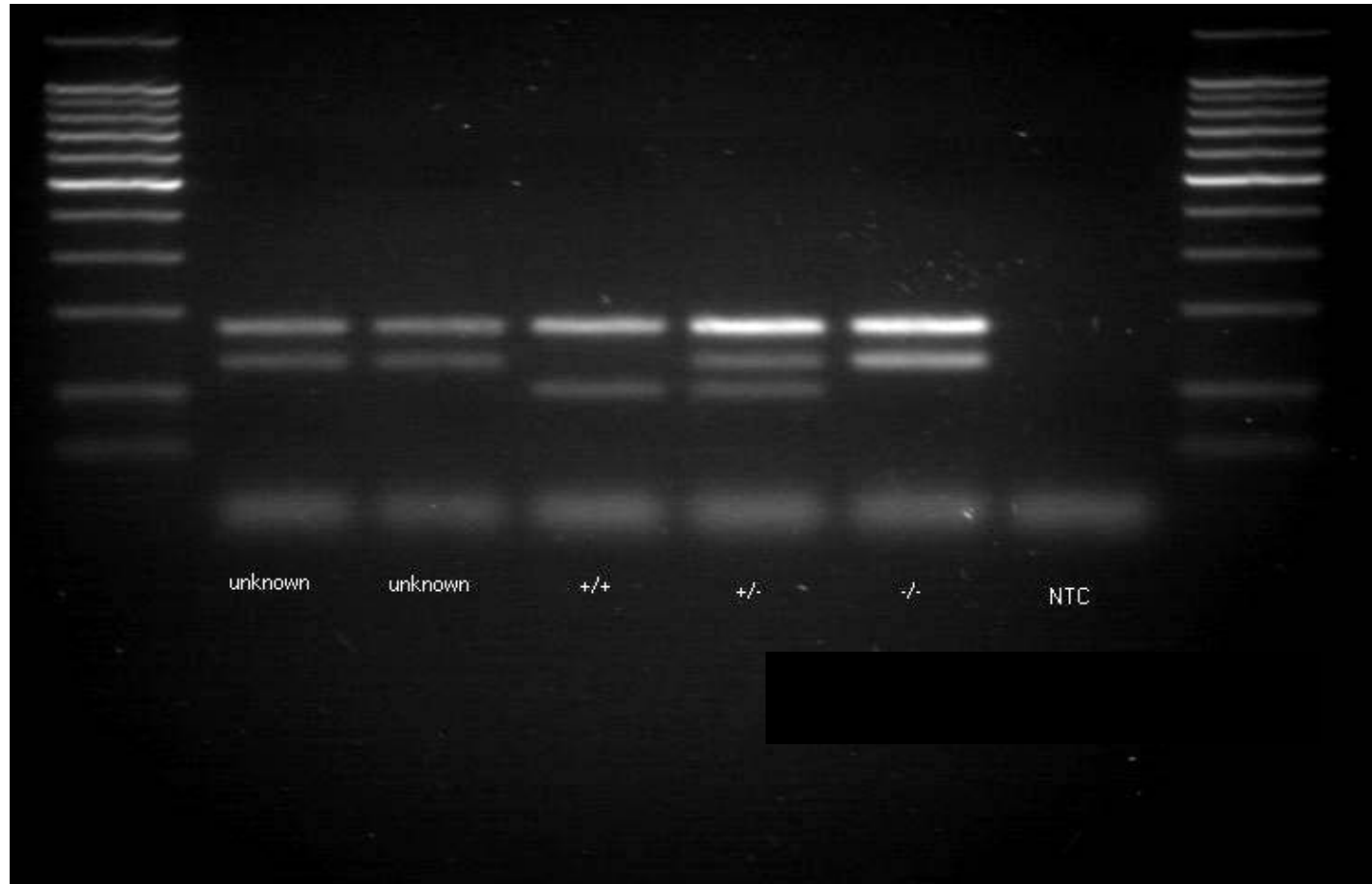
- **Neonates with Classic MSUD**

- maple syrup odor (12-24 HR)
- irritability, and poor feeding (2-3 days)
- lethargy, intermittent apnea, opisthotonus, "fencing" and "bicycling" (4-5 days)
- central respiratory failure (7-10 days)

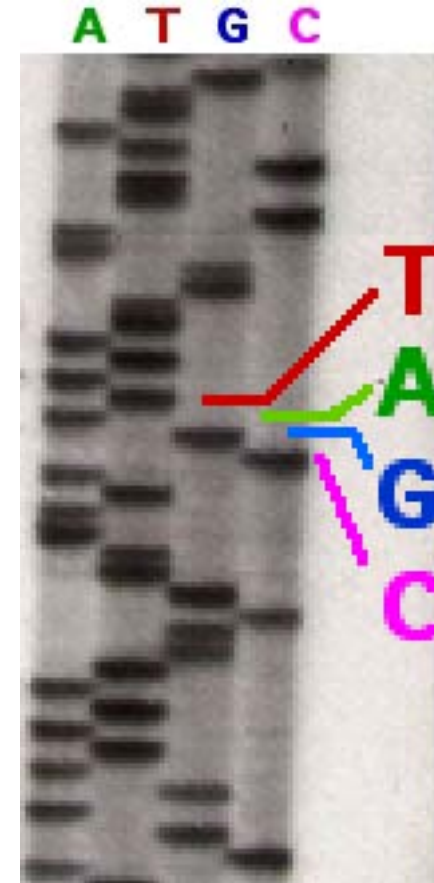
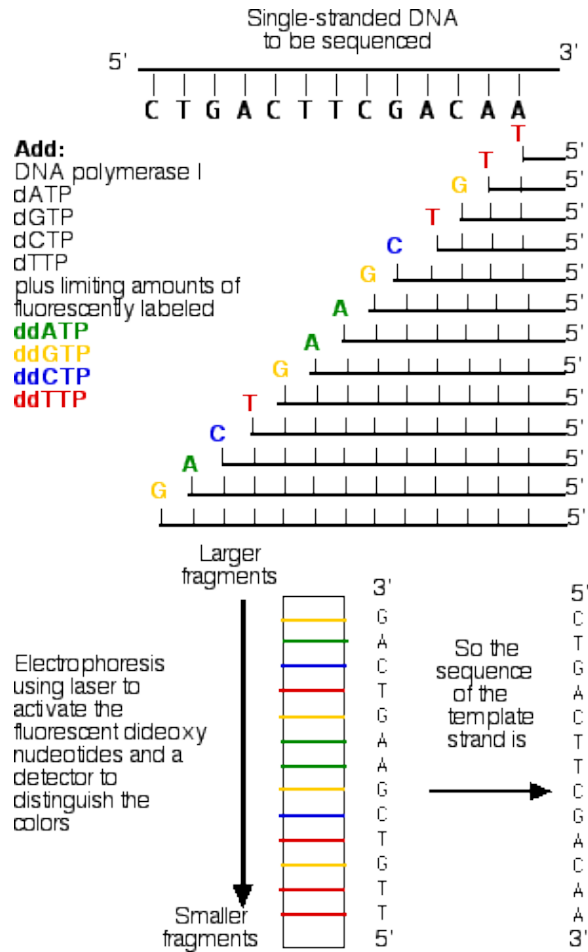
- **MSUD in Old Order Mennonites**

- Carrier frequency is as high as one in ten
- Disease incidence of approximately one in 380 live births
- Founder [mutation](#) (c.1312T>A) in *BCKDHA*

***BCKDHA* c. 1312T>A Mutation detection using Tetra-primer ARMS-PCR**

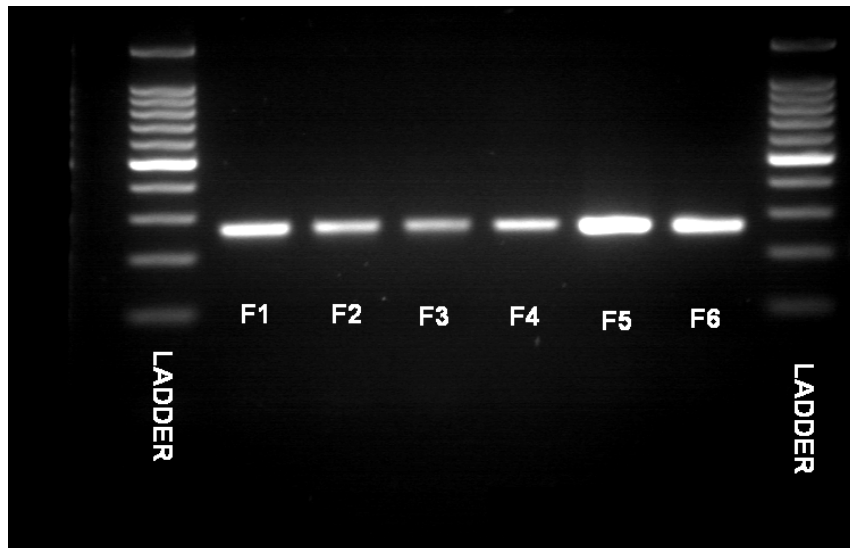


Sanger's Method of DNA Sequencing

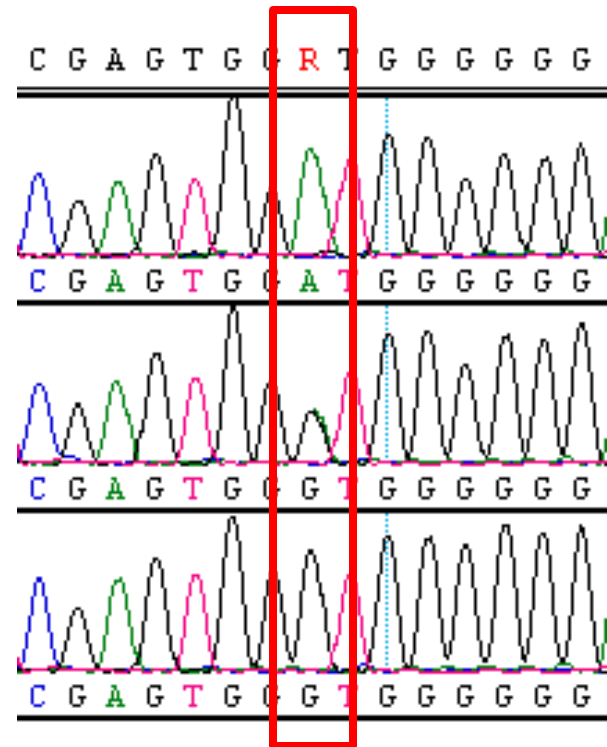


ACADSB c.1165A>G Mutation Detected by Sanger Sequencing

PCR products flank *ACADSB* c.1165A>G site

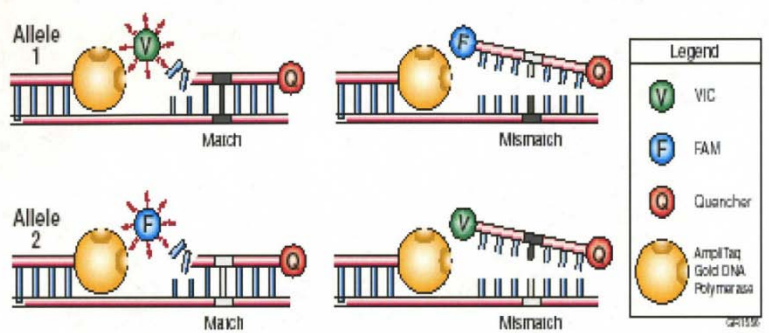


Sequencing Results



Greg Kopish & Timothy Davis

Allelic Discrimination Using TaqMan Probes



The diagram shows four scenarios of TaqMan probe discrimination:

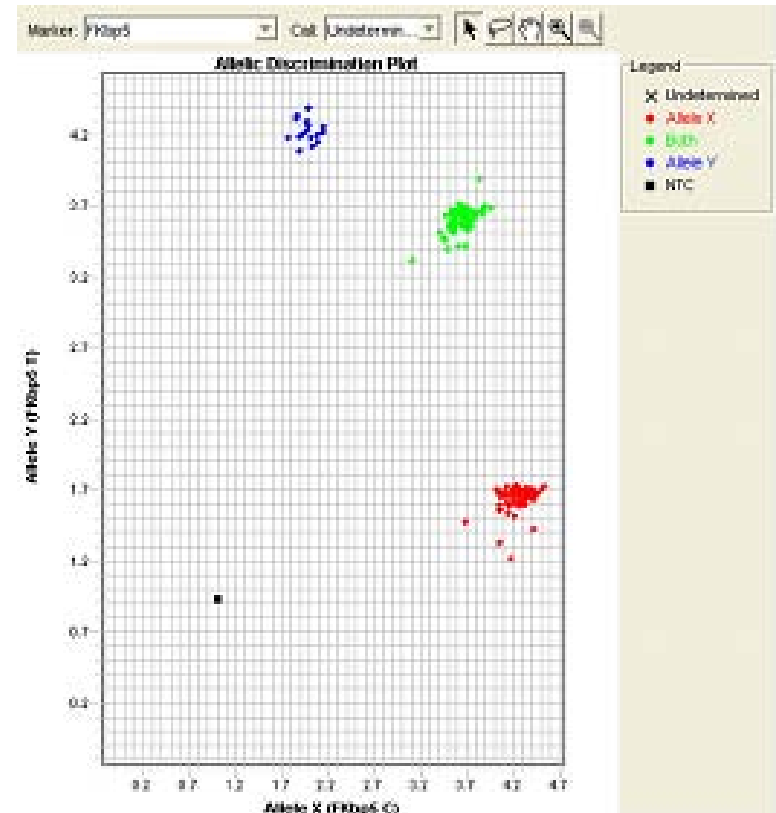
- Allele 1 Match:** The probe hybridizes perfectly, and the polymerase cleaves it, releasing the VIC fluorophore (green).
- Allele 1 Mismatch:** The probe has a mismatch, preventing cleavage and releasing the FAM fluorophore (blue).
- Allele 2 Match:** The probe hybridizes perfectly, and the polymerase cleaves it, releasing the FAM fluorophore (blue).
- Allele 2 Mismatch:** The probe has a mismatch, preventing cleavage and releasing the VIC fluorophore (green).

Legend:

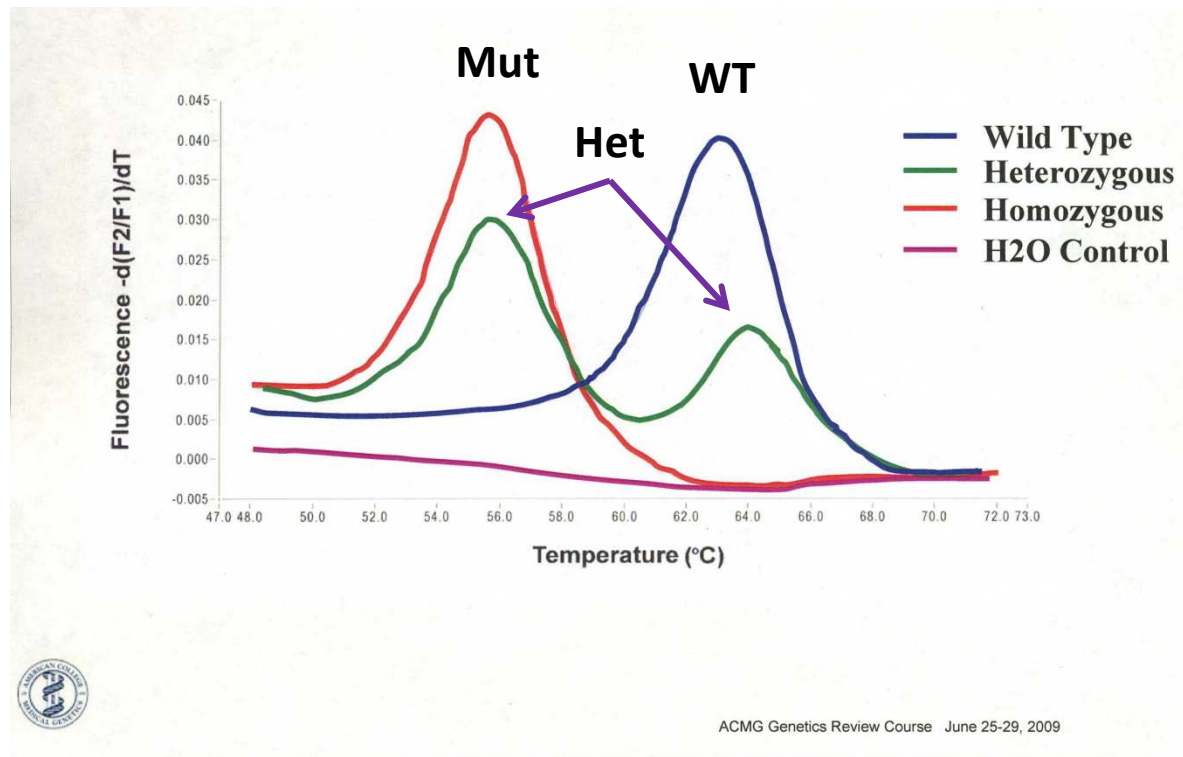
- VIC (Green circle)
- FAM (Blue circle)
- Quencher (Red circle)
- AmpTaq Gold DNA Polymerase (Orange circle)

The table below summarizes the possible results of the example allelic discrimination assay shown above.

A substantial increase in...	Indicates...
VIC fluorescence only	homozygosity for Allele 1.
FAM fluorescence only	homozygosity for Allele 2.
both fluorescent signals	heterozygosity.



Fluorescence Resonance Energy Transfer (FRET) Real-time PCR Assay



Summary

Assay	Procedures	Instrument
RFLP	Conventional PCR Restriction enzyme digestion Agarose gel electrophoresis (Size-based discrimination)	Thermal cycler Gel electrophoresis unit
ARMS-PCR	Conventional PCR Agarose gel electrophoresis (Size-based discrimination)	Thermal cycler Gel electrophoresis unit
Targeted gene sequencing	Conventional PCR PCR products treatment Sequencing Reaction Capillary electrophoresis	Thermal cycler DNA analyzer
TaqMan real-time PCR	Real-time PCR	Real-time PCR system
FRET real-time PCR	Endpoint PCR	Real-time PCR system

**Success is the sum of a lot of small things
correctly done.**

Chef Fermand Point