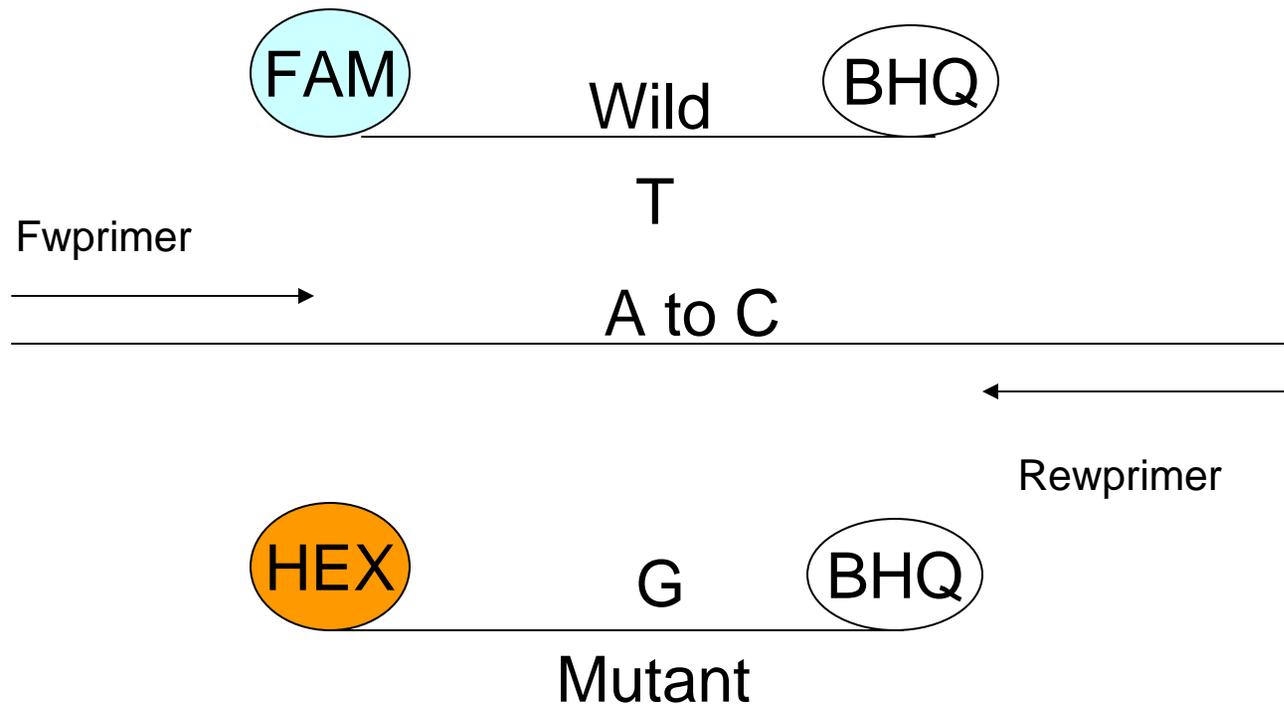




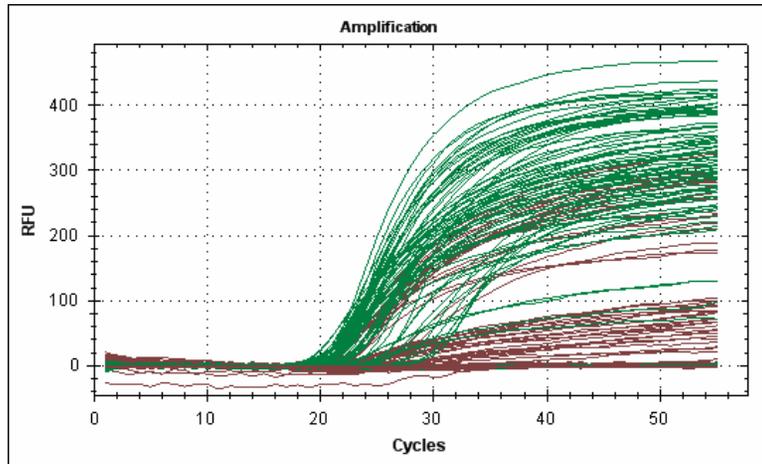
## SNP-detection with Allele-Specific Probes

Concept:



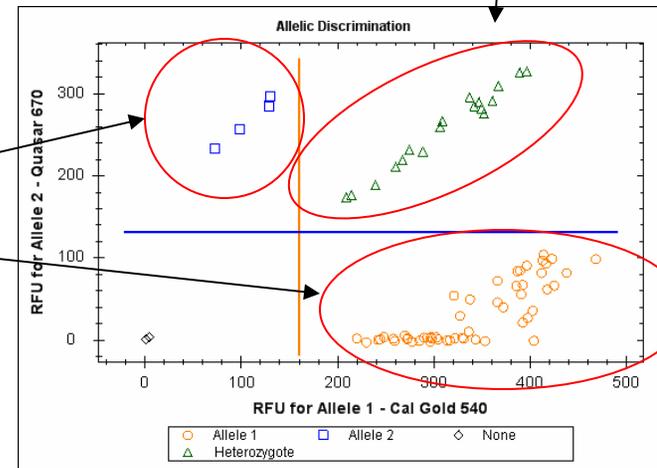


# Allelic discrimination module



Homozygotes

Heterozygotes





# Allelic discrimination module

Bio-Rad CFX Manager

Data Analysis - 2C192 20081117 1.pcrd

File View Settings Tools

Well Group: All Wells

Detected Instruments

Quantitation Quantitation Data Gene Expression End Point Allelic Discrimination QC Run Information

**Allelic Discrimination**

Well	Sample	RFU1	RFU2	Call	Type
A01	2C19*2/HH069	294	3.12	Allele 1	Auto
A02	2C19*2/HH069	298	2.06	Allele 1	Auto
A03	2C19*2/HH073	322	2.46	Allele 1	Auto
A04	2C19*2/HH073	303	3.43	Allele 1	Auto
A05	2C19*2/HH074	130	296	Allele 2	Auto
A06	2C19*2/HH074	129	285	Allele 2	Auto
A07	2C19*2/HH076	72.0	233	Allele 2	Auto
A08	2C19*2/HH076	98.0	256	Allele 2	Auto
A09	2C19*2/HH081	317	-0.0112	Allele 1	Auto
A10	2C19*2/HH081	314	-0.179	Allele 1	Auto
A11	2C19*2/HH084	285	-0.698	Allele 1	Auto
A12	2C19*2/HH084	304	1.03	Allele 1	Auto
B01	2C19*2/HH085	273	2.46	Allele 1	Auto
B02	2C19*2/HH085	290	3.26	Allele 1	Auto
B03	2C19*2/HH086	390	84.7	Allele 1	Auto
B04	2C19*2/HH086	417	93.0	Allele 1	Auto

	1	2	3	4	5	6	7	8	9	10	11	12
A	Unk											
B	Unk											
C	Unk											
D	Unk											
E	Unk											
F	Unk											
G	Unk	Unk	Unk	NTC								
H												

Selected Fluorophores (X = Allele1, Y = Allele2)  
 X: Cal Gold 540 Y: Quasar 670  
 Call Selected Alleles: Automatic  
 Display Mode:  Ct  RFU Select Cycle: 55  Normalize Data  
 Thresholds: Vertical: 160.185 Horizontal: 131.307  
 Restore Default Thresholds

Completed Scan Mode: All Channels Plate Type: BR Clear Analysis Mode: Baseline Subtracted Curve Fit

No instrument detected after 120 seconds.

User:admin 5/16/2010 5:02 PM

start Jozsef Paulik - Inbox ... MEA\_Roadshow Microsoft PowerPoint ... Bio-Rad CFX Manager ... Data Analysis - 2C19 ... 5:02 PM