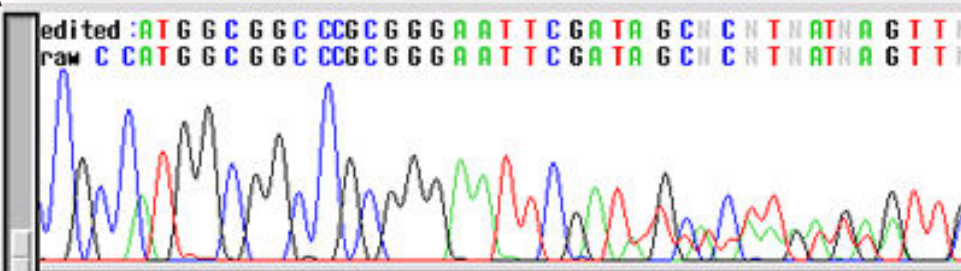


Molecular Markers



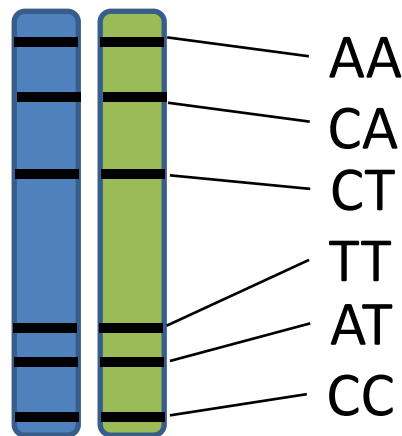
CRITFC Genetics Workshop
December 9, 2014

Molecular Markers

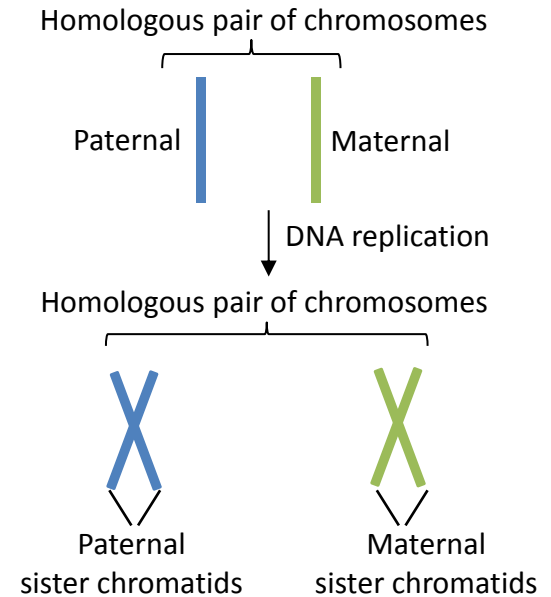
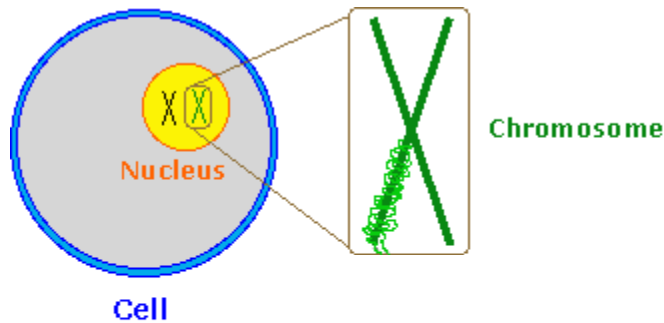
- Tools that allow us to collect information about an individual, a population, or a species
- Application in fisheries
 - mating behaviors/effective population size
 - population structure
 - parentage, genetic tagging
 - genetic basis of traits/gene mapping

Molecular Markers

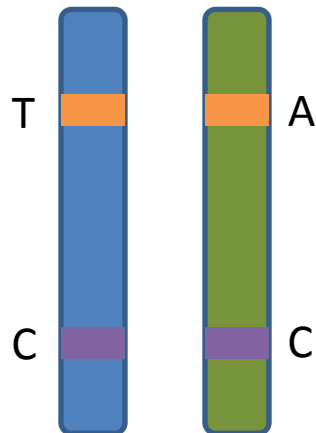
- Designed to interrogate heritable differences in DNA sequence called “polymorphisms”
- Individuals may inherit unique combinations of polymorphisms/alleles across several loci providing a distinctive DNA ID or fingerprint



Molecular Markers



Homologous pair of chromosomes



Locus – physical location within the DNA sequence

Allele – a single variant of a locus

Genotype – a set of 2 alleles at a locus

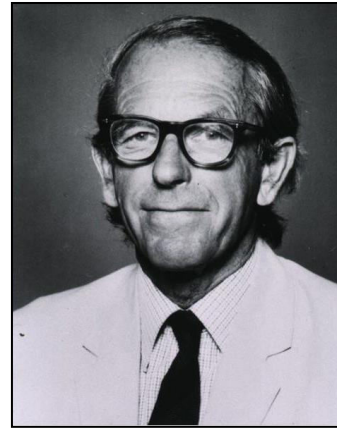
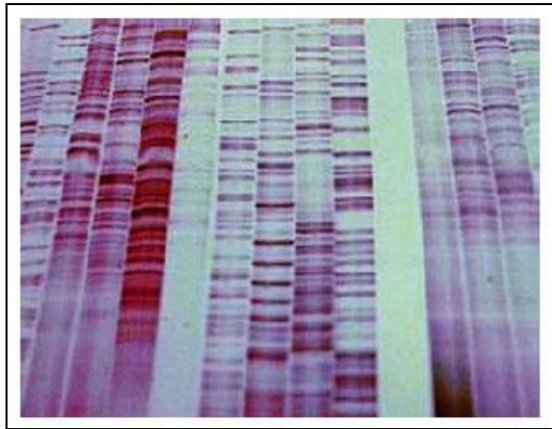
2 loci:

TA = heterozygous

CC = homozygous

DNA Sequence and Genetic Markers

- 1977 Sanger and colleagues describe laboratory methods of DNA sequencing

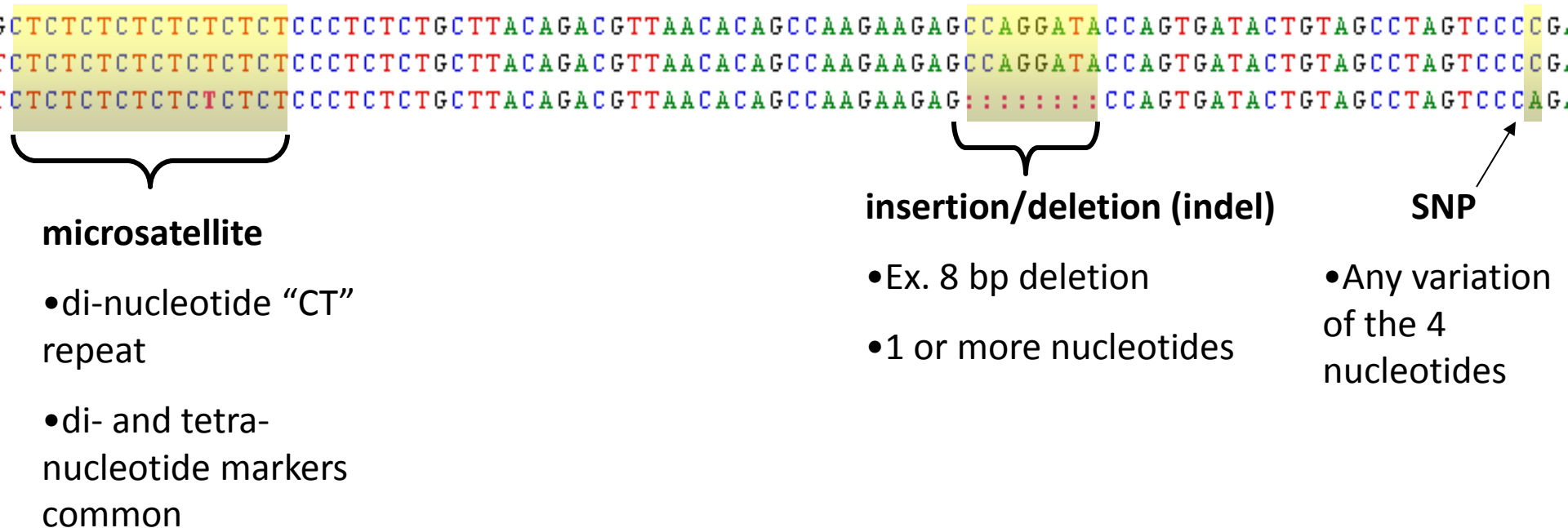


- Allowed for the direct interrogation of variation at the DNA sequence level which lead to the discovery of informative genetic markers

DNA Sequence Variation

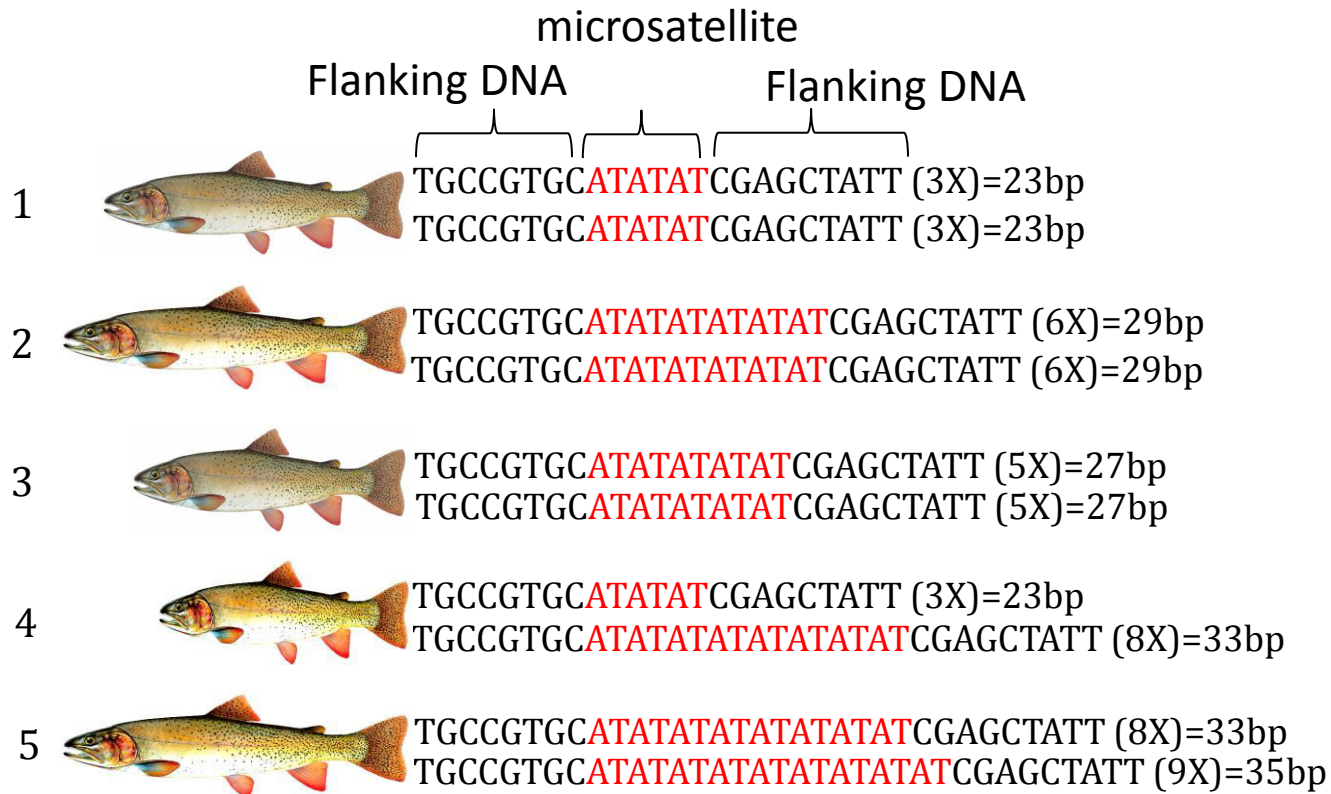
Mistakes made during DNA replication leads to sequence variation (i.e., mutations)

DNA sequence of three individuals:



Microsatellite Markers

Repeating sequences of 2-6 base pairs of DNA
and can be hyper-variable compared to other markers

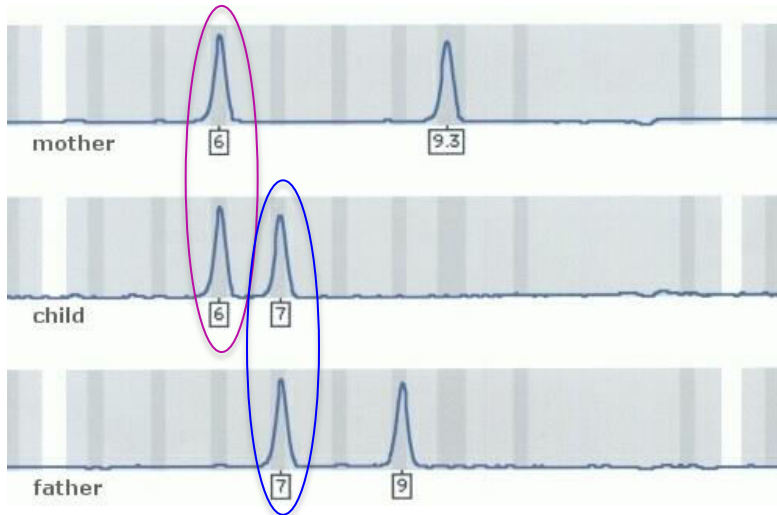


Flanking DNA is identical, but length of microsatellite is different between individuals

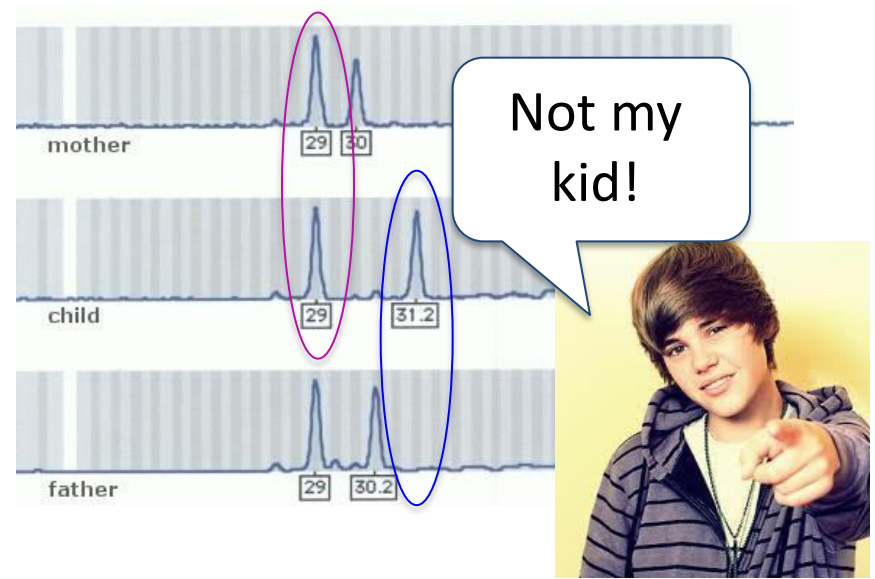
Microsatellite Markers

- Example: Paternity testing

Case 1



Case 2



Microsatellite Markers

- Pros
 - Extremely variable
 - > 20 alleles for one locus
 - Moderately abundant in genome
 - Predominantly neutral loci
- Cons
 - Difficult to discover
 - Generally requires DNA sequence information
 - Difficult to standardize and exchange data across labs
 - PCR variation
 - Scoring variation

SNP Markers

Single nucleotide polymorphism

<https://www.youtube.com/watch?v=tJjXpiWKMyA>

- Variation in DNA, when a single nucleotide (A, T, C, or G) within a given sequence differs between homologous chromosomes or between individuals at homologous loci



ATG	GCT	TCG	ATC	GAT	CTA
ATG	GCC	TCG	ATC	GAT	CTA



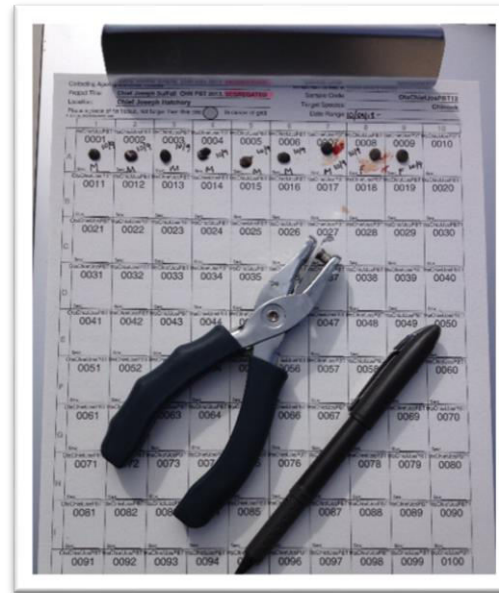
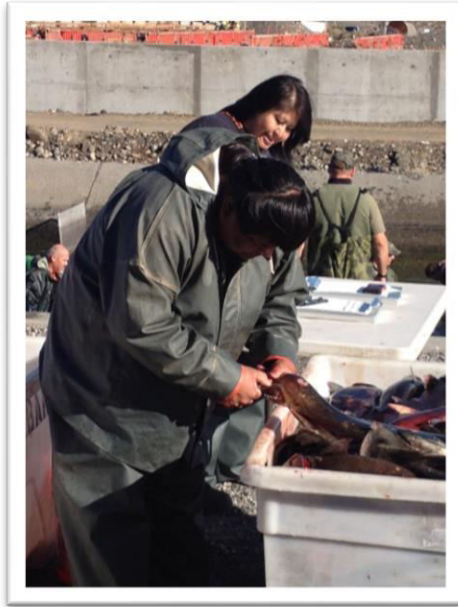
ATG	GCT	ACG	ATC	GAC	CTA
ATG	GCT	ACG	ATC	GAC	CTA

SNP Markers

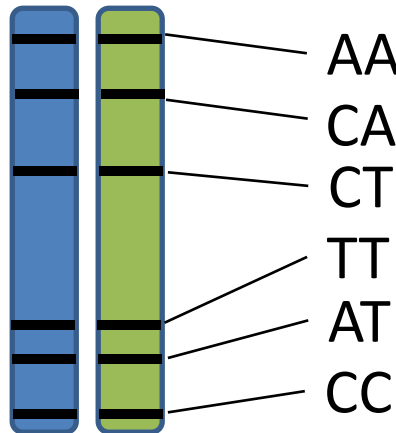
- Pros
 - Codominant markers
 - Most abundant markers in the genome
 - Easy to interrogate with current high-throughput technology
 - requires little tissue
 - Highly reproducible between labs, easy to standardize, easy exchange of data
 - Can be adaptive and neutral loci
- Cons
 - Not as variable as microsatellites
 - High up front discovery/operating cost

Questions?

Laboratory techniques

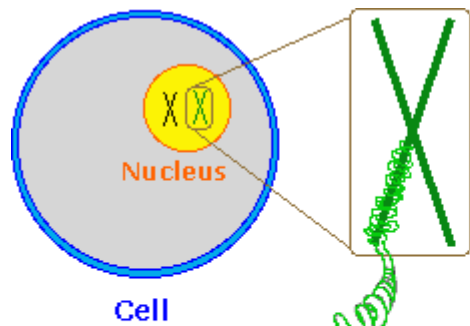


How do we get from tissue sample
to multi-locus genotype?



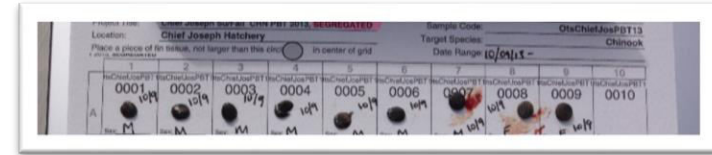
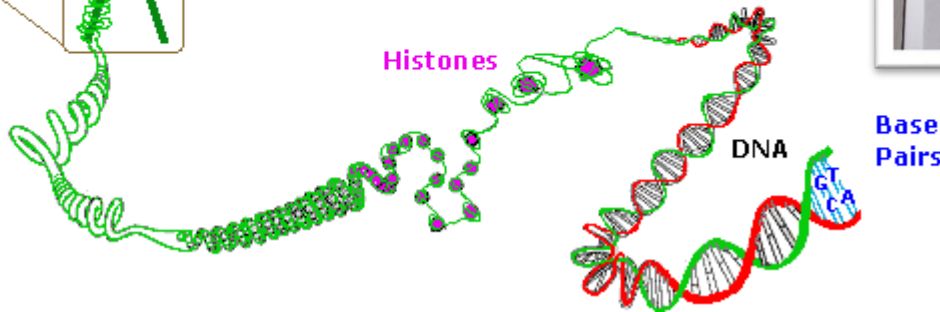
1.) DNA extraction

2.) Amplify specific regions of
DNA (i.e., molecular markers)
via PCR



Chromosome

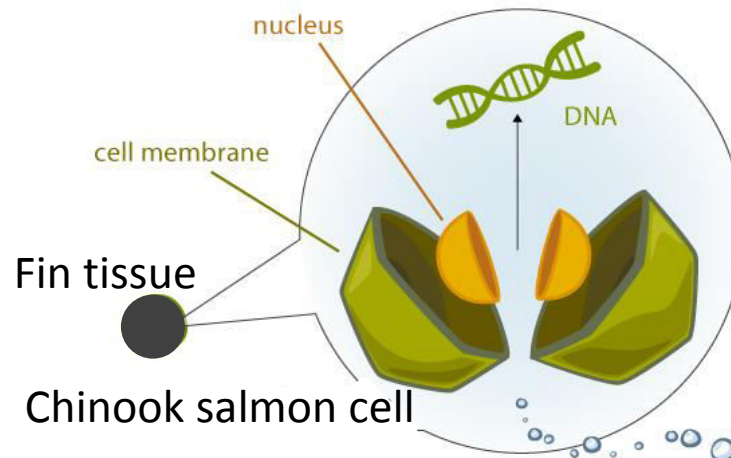
DNA Extraction

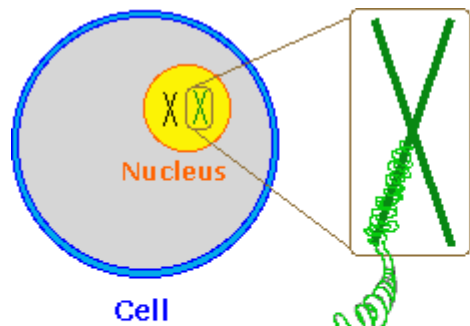


Ingredients

- Tissue
- Detergent & Salt
- Protease
- Alcohol

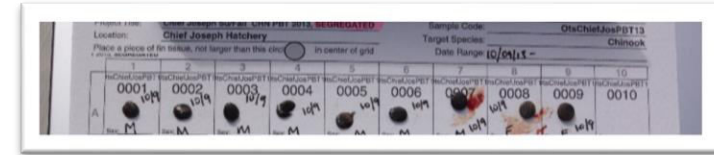
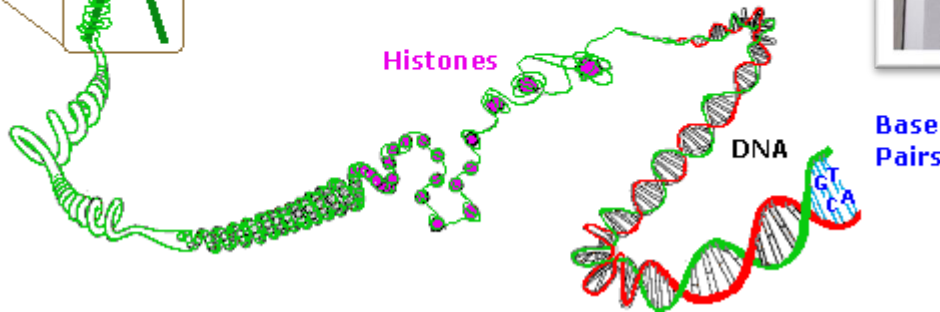
Break up cell walls, disrupt proteins





Chromosome

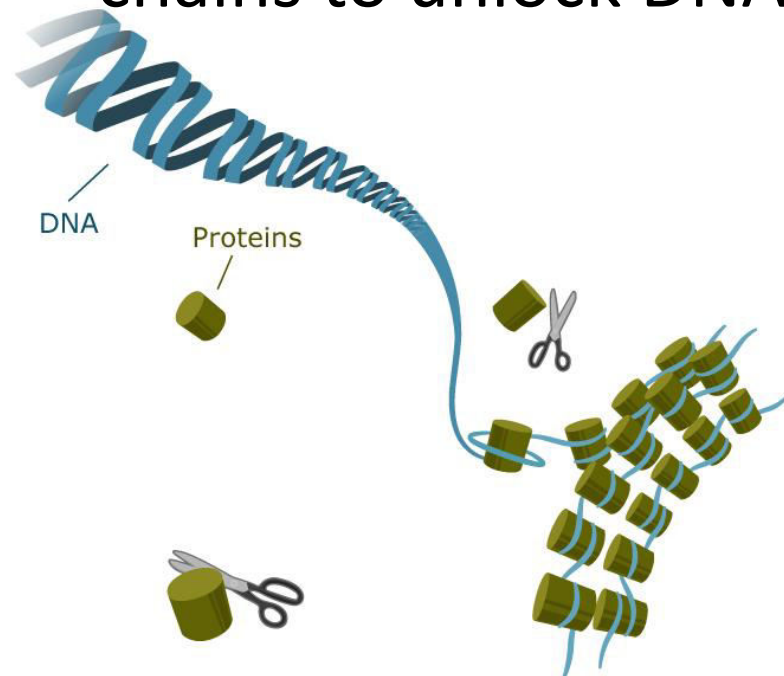
DNA Extraction

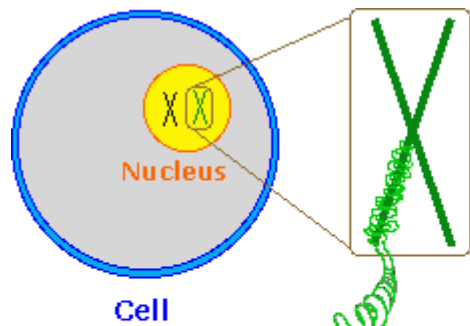


Ingredients

- Tissue
- Detergent & Salt
- Protease
- Alcohol

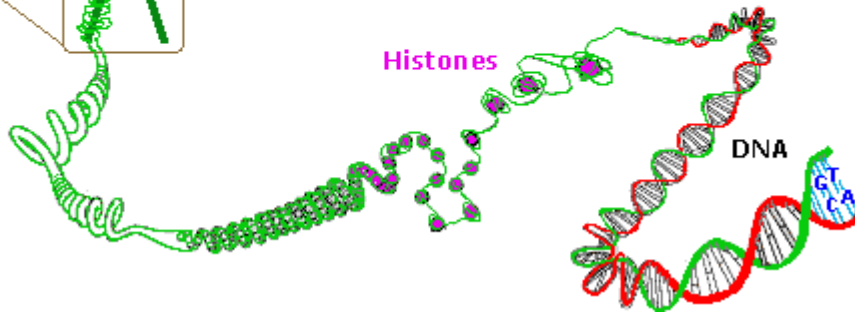
Emzyme cuts amino acid chains to unlock DNA





Chromosome

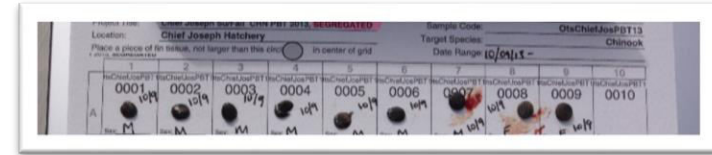
DNA Extraction



Histones

DNA

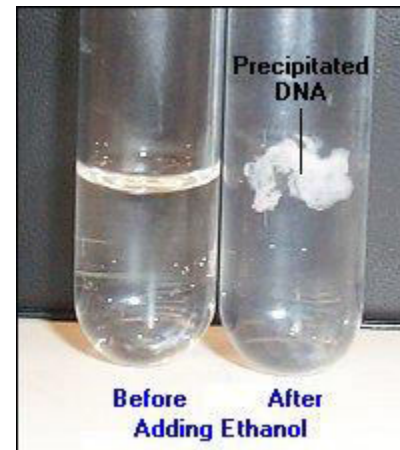
Base Pairs



Ingredients

- Tissue
- Detergent & Salt
- Protease
- Alcohol

DNA is not soluble in ethanol, and will precipitate



DNA Extraction

Qiagen Dneasy kits

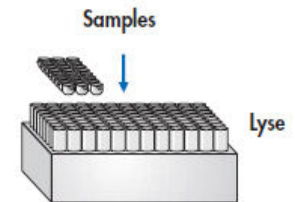


- Small piece of fin clip is added to lysis buffer
- Tissue is digested

DNeasy Mini Procedure



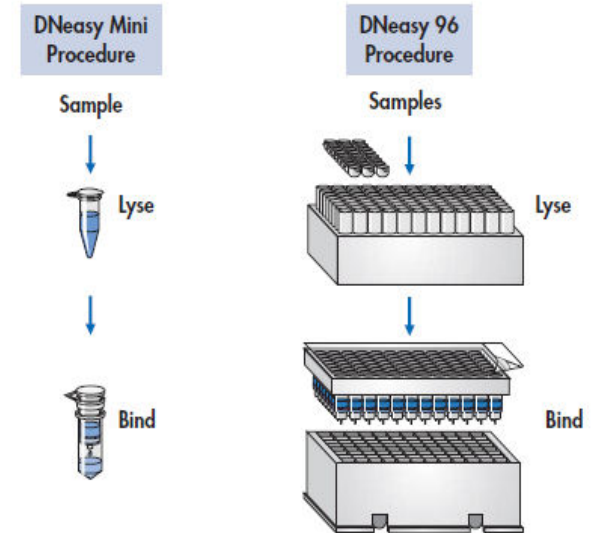
DNeasy 96 Procedure



DNA Extraction Qiagen Dneasy kits



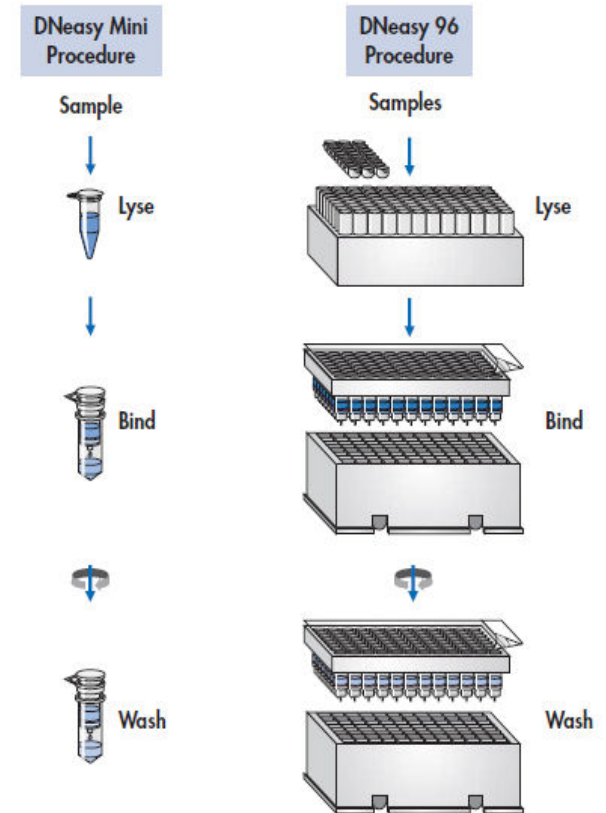
- Small piece of fin clip is added to lysis buffer
- Tissue is digested
- Add solution containing alcohol
- Precipitated DNA is caught in filter columns



DNA Extraction Qiagen Dneasy kits



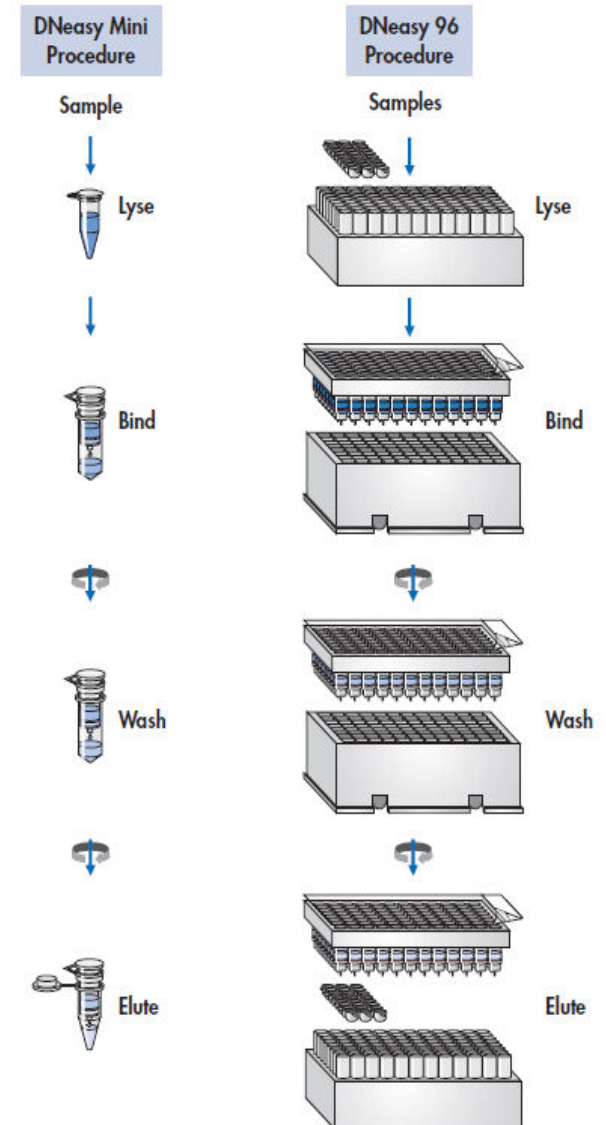
- Small piece of fin clip is added to lysis buffer
- Tissue is digested
- Add solution containing alcohol
- Precipitated DNA is caught in filter columns
- 2 wash steps to remove salts



DNA Extraction Qiagen Dneasy kits



- Small piece of fin clip is added to lysis buffer
- Tissue is digested
- Add solution containing alcohol
- Precipitated DNA is caught in filter columns
- 2 wash steps to remove salts
- DNA is released from filter by dissolving in weak buffer



PCR: Polymerase Chain Reaction

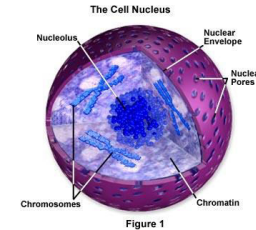
Review what happens in the cell (DNA replication): <http://www.dnalc.org/resources/3d/03-mechanism-of-replication-basic.html>

Generates millions of copies of a particular sequence (i.e., microsatellite, SNP) of DNA



PCR ingredients

- Genomic DNA
- Primers
- Taq polymerase
- Nucleotides (A, C, T, G)



Cell ingredients

- Genomic DNA
- Primase enzyme
- DNA polymerase
- Nucleotides (A, C, T, G)

PCR: Polymerase Chain Reaction

Amplifying a microsatellite



pcr.exe

Tell DNA from which you will amplify a
Synthesizes DNA by adding nucleotides
molecular marker

ATGCAGGCTGA**GAGAG**ACTAGTCGATG
CAGCTAC

ATGCAGGCTGA
TACGTCCGACT**CTCTCT**GATCAGCTAC

Ingredients

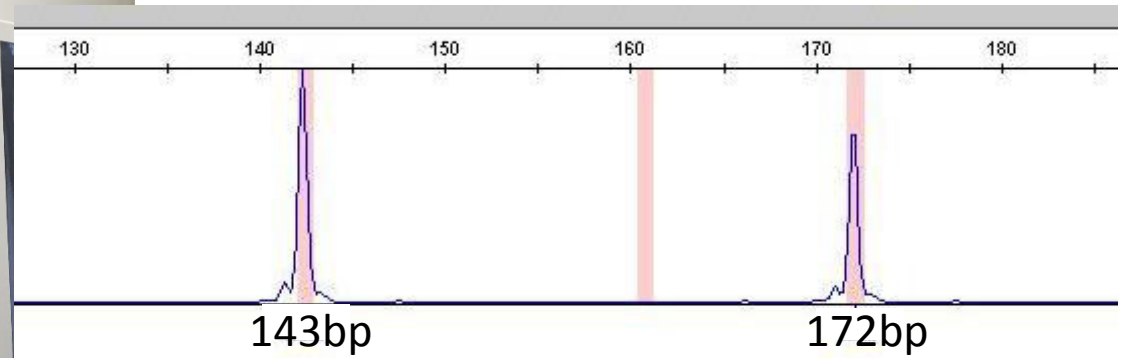
- Genomic DNA
- Primers
- Enzyme (Taq)
- Nucleotides (A, C, T, G)



PCR: Polymerase Chain Reaction

“Reading” a microsatellite

Sizes of PCR products determined by the 3730 DNA analyzer



	A	B	C	D	E	F	G	H	I	J	K
1	Individual Name	usat1a	usat1b	usat2a	usat2b	usat3a	usat3b	usat4a	usat4b	usat5a	usat5b
2	Sample 1	142	154	241	243	132	144	142	142	241	243
3	Sample 2	164	164	251	251	144	144	142	142	243	251
4	Sample 3	142	164	243	243	132	132	142	142	243	243
5	Sample 4	154	164	241	243	132	144	142	152	241	243
6	Sample 5	154	154	253	243	154	154	142	142	253	243
7	Sample 6	154	176	241	243	132	154	142	142	243	243

PCR: Polymerase Chain Reaction

Amplifying a SNP

<http://www.lifetechnologies.com/us/en/home/life-science/pcr/real-time-pcr/qpcr-education/ask-taqman/ask-taqman-video-gallery.html>

Ingredients

- Genomic DNA
- Primers
- Enzyme (Taq)
- Nucleotides (A, C, T, G)
- Fluorescent probes

ATGCAGGCTGA^GCCATGCTAGTCGATG
CAGCTAC

ATGCAGG
TACGTCCGACT^CGGTACGATCAGCTAC

●AGCGA●

●ATCGA●

PCR: Polymerase Chain Reaction

Amplifying a SNP

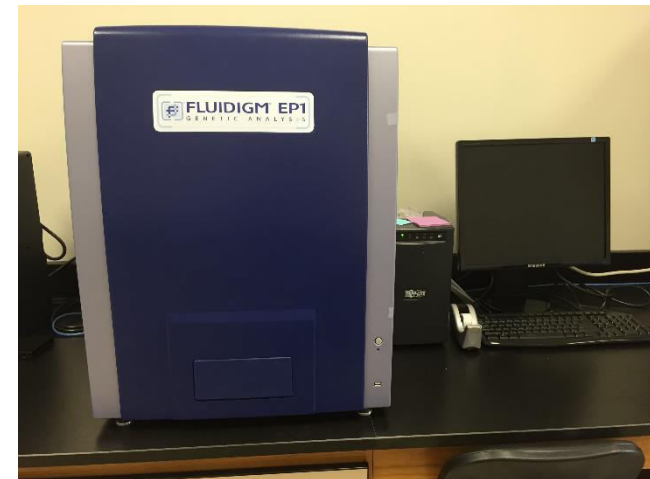
Ingredients

- Genomic DNA
- Primers
- Enzyme (Taq)
- Nucleotides (A, C, T, G)
- Fluorescent probes

ATGCAGGCTGA[●]CCATGCTAGTCGATG
CAGCTAC

ATGCAGG[●]
TACGTCCGACT[●]CGGTACGATCAGCTAC
AGCGA[●]

ATCGA[●]

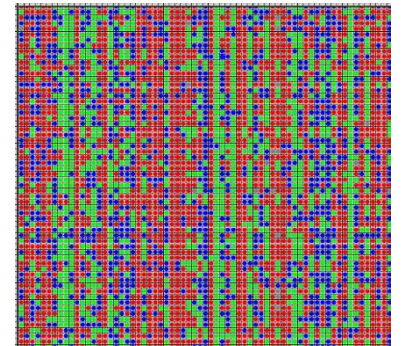
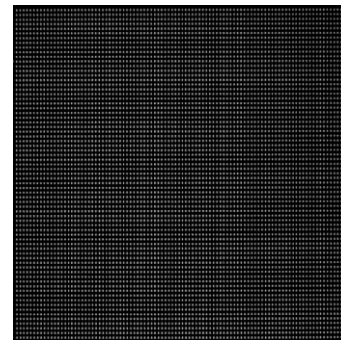
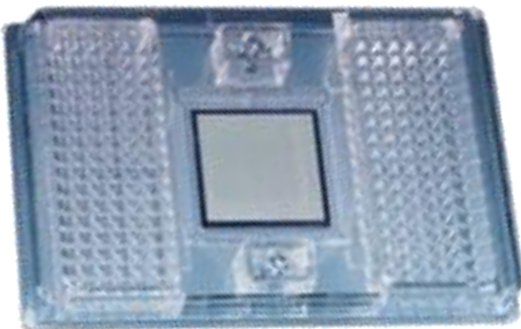


PCR: Polymerase Chain Reaction

Amplifying a SNP

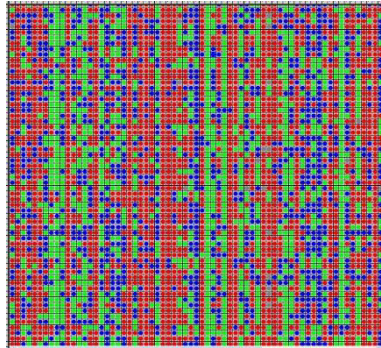
General Steps

- Pre-amplify DNA (boost starting copy; 14 cycles)
- 1 chip = 96 SNP assays, 96 samples
- Array is loaded with PCR reagents and the pre-amplified samples
- PCR resumes inside the chip for 50 cycles in the FC-1 thermal cycler
- Chip is then read using the camera inside the EP-1 instrument.

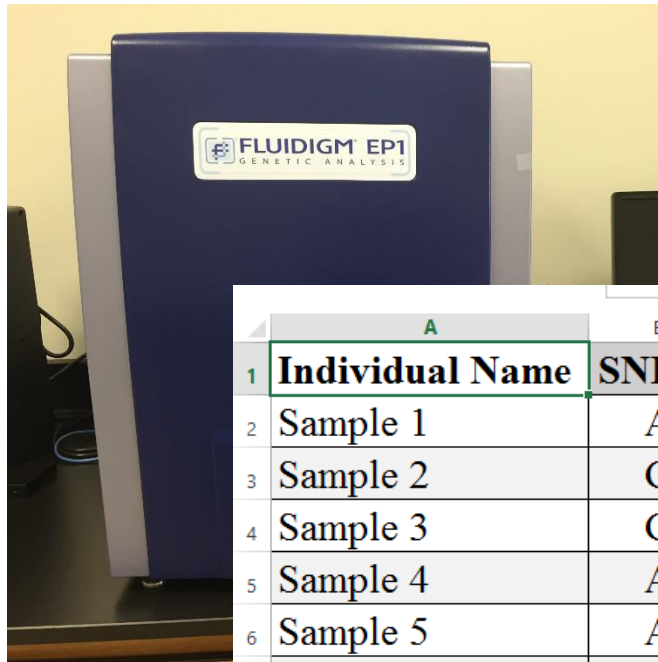


PCR: Polymerase Chain Reaction

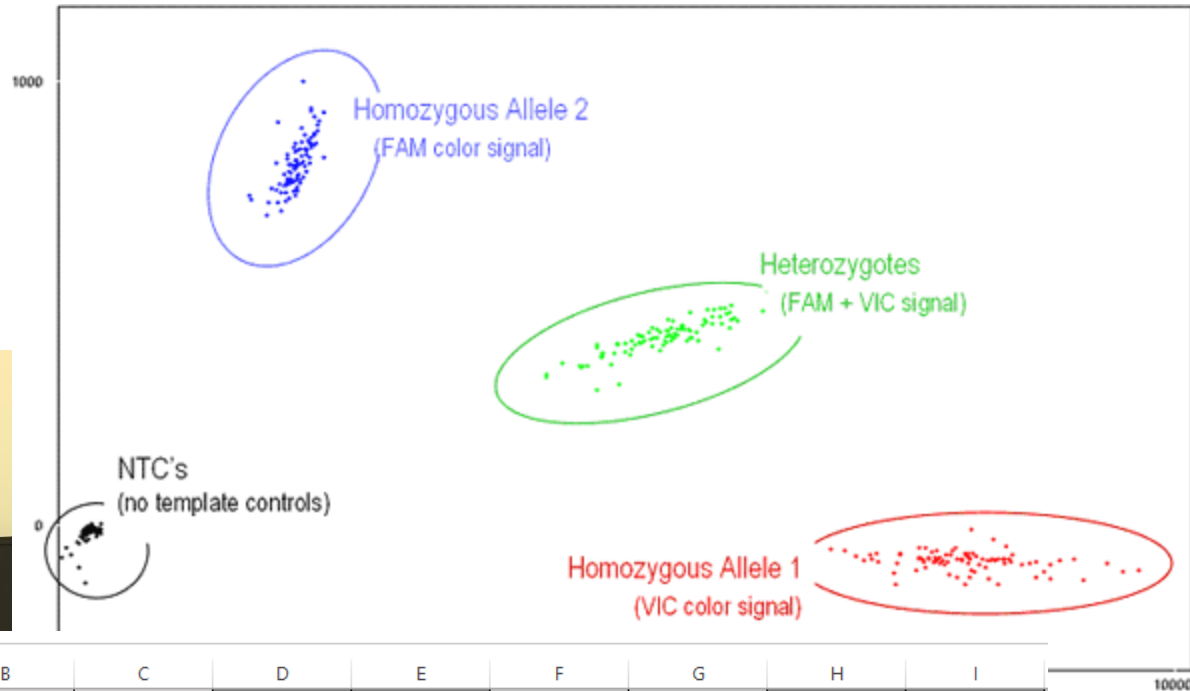
“Reading” a SNP



Fancy
machine
reads the
chip



Example plot – 96 samples for 1 SNP



	A	B	C	D	E	F	G	H	I
1	Individual Name	SNP1a	SNP1b	SNP2a	SNP2b	SNP3a	SNP3b	SNP4a	SNP4b
2	Sample 1	A	A	C	G	C	C	C	C
3	Sample 2	G	G	G	G	T	T	C	C
4	Sample 3	G	G	C	C	C	T	C	C
5	Sample 4	A	A	G	G	C	T	C	C
6	Sample 5	A	G	C	C	C	T	C	C
7	Sample 6	A	G	C	G	C	C	C	C

SNPs and next generation sequencing

“Reading DNA” – determine order of the 4 nucleotides

<http://www.dnalc.org/view/15479-Sanger-method-of-DNA-sequencing-3D-animation-with-narration.html>



- Genome sequencing and assembly
- RNA-seq: mRNA transcript analysis (differences in gene expression)
- RAD-seq: method to discover and genotype ~10K SNP markers per 50-100 samples
- GT-seq: genotype by sequencing; ~2,000 samples in lane, ~192 SNPs