

# Genetic test for identification of parents (parentage testing)

## Lecture

**Modul no. 1: Precision livestock farming**

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# Basic terms

parentage: parenthood

paternity: fatherhood

Individual identification: determining that the DNA sample belongs to a specific individual (molecular fingerprinting)

Genetic type: a set of genotypes (alleles) characteristic and unmistakable for everyone, also known as *genetic profile*



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# Parentage testing and individual identification in animals

- importance - verification of origin and identification
- livestock in CZ (it varies by country)
  - pig: determining of the genetic type (identification) of boars in breeding farms
  - cattle: genetic type before selection for breeding (bull breeding), verification and certification of the origin of breeding bulls
  - horse: verification and certification of the origin of foals after insemination (always for English thoroughbred, even if there is no insemination)
  - other species: dog, exotic birds, protected animals CITES (origin and identification)



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# Importance

**Importance** in animal genetics, forensic genetics and medicine

## *Animals:*

- frequent errors in records

- incorrect parentage data to estimate breeding value

- you buy an animal that does not belong to the declared (quality) parents

## *Human:*

- paternity disputes

- identification of the offender, or victims

- transplantation and other medical procedures where kinship is decisive



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# Development and use of methods

Previously: blood group polymorphism and protein

New: DNA variability (DNA fingerprinting)

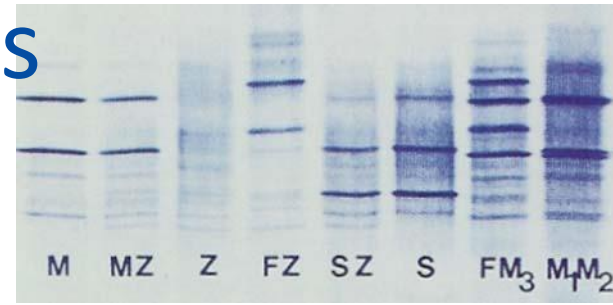
- VNTR (variable number of tandem repeats, minisatellites)

Currently:

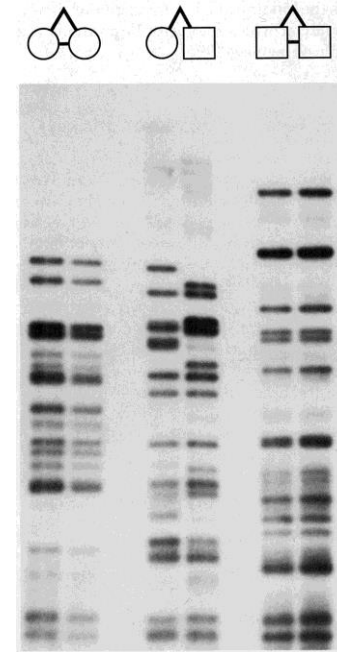
- STR (short tandem repeats, microsatellites)
- SNP (latest, increasingly used)

## Requirements:

- high reliability of rejection and (or) confirmation of parentage
- relatively easy and quick laboratory testing
- high reproducibility
- the lowest price



*protein IEF*



*VNTR*



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# Microsatellites

- short tandem repetitions composed of mono, di, tri or tetra nucleotide motifs
- highly polymorphous  $\Rightarrow$  different alleles (5-20)

e.g.  $(GC)_n$   $n = 5-25$  repetitions

- length polymorphism - i.e. if the MS sequence is amplified by PCR, we detect by electrophoresis a different fragment size (allele) due to a different number of repetitions
- the allele designation is therefore the base pair size

# Mikrosatelites

↓ primer forward

mikrosatelite

primer reverse ↓

(GC)<sub>n</sub>  
PCR size

5' **CATTGAATCGGTATCAT**(N)<sub>60</sub>**AGCGCGCGCGCGCGCGCT**(N)<sub>40</sub>**ACGTTAATTCATTTCGTG** 3' n=8  
152 bp

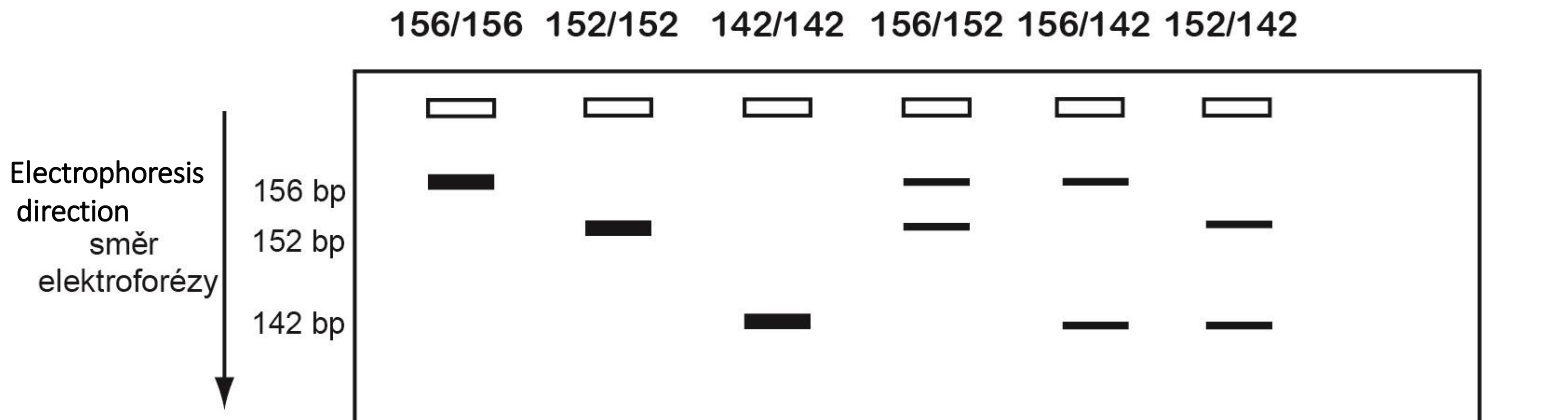
5' **CATTGAATCGGTATCAT**(N)<sub>60</sub>**AGCGCGCT**(N)<sub>40</sub>**ACGTTAATTCATTTCGTG** 3' n=3  
142 bp

5' **CATTGAATCGGTATCAT**(N)<sub>60</sub>**AGCGCGCGCGCGCGCGCGCT**(N)<sub>40</sub>**ACGTTAATTCATTTCGTG** 3' n=10  
156 bp

PAGE scheme

Schéma PAGE

genotype



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# Used microsatellites

Examples:

*Pig* (10x) – S0068, S0107, SW24, SW840, SW353, SW936, SW353, S0070, SW72, TNFB

*Cattle* (11x) – TGLA53, TGLA122, TGLA126, TGLA227, ETH3, ETH10, ETH225, BM1824, BM2113, INRA23, SPS115

*Horse* (17x) – VHL20, AHT4, HTG4, HMS7, HTG6, AHT5, HMS6, ASB23, ASB2, HTG10, HTG7, HMS2, HMS3, ASB17, LEX3, HMS1, CA425

*Dog* (18x) – AHTk211, CXX279, INU055, REN169018, REN54P11, AHT137, AHTTh260, AHTk253, INRA21, REN169D01, AHT121, AHTTh171, FH2054, REN162C04, REN247M23, FH2848, INU005, INU30



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# Additional panels

- raptor 5 MS
- goat 14 MS
- sheep 13 MS
- deer, cats, beavers, nutria, hens, geese, llamas



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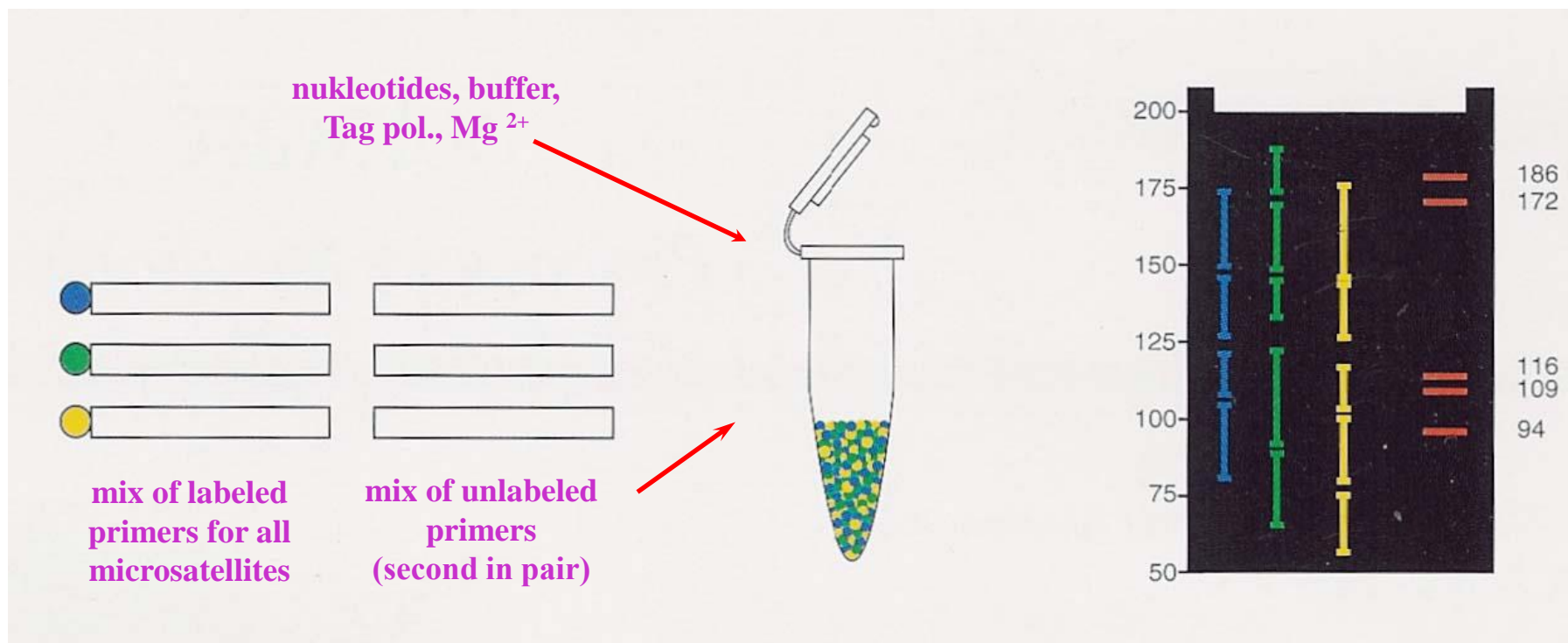
# Analysis procedure

1. Isolation of the genomic DNA of the tested individual
2. Multiplex PCR - amplification of a panel of microsatellites
3. Fragmentation analysis using fluorescence capillary electrophoresis
4. Determination of peak sizes
5. Determination of alleles of individual microsatellites: discrimination by color and known range of alleles
6. Evaluation - verification of parentage



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# Multiplex PCR



the allele size ranges of different microsatellites in the same color must not overlap

Fluorescent colours(G5):

FAM - blue

VIC - green

NED – yellow

PET - red

LIZ – orange (size standard)

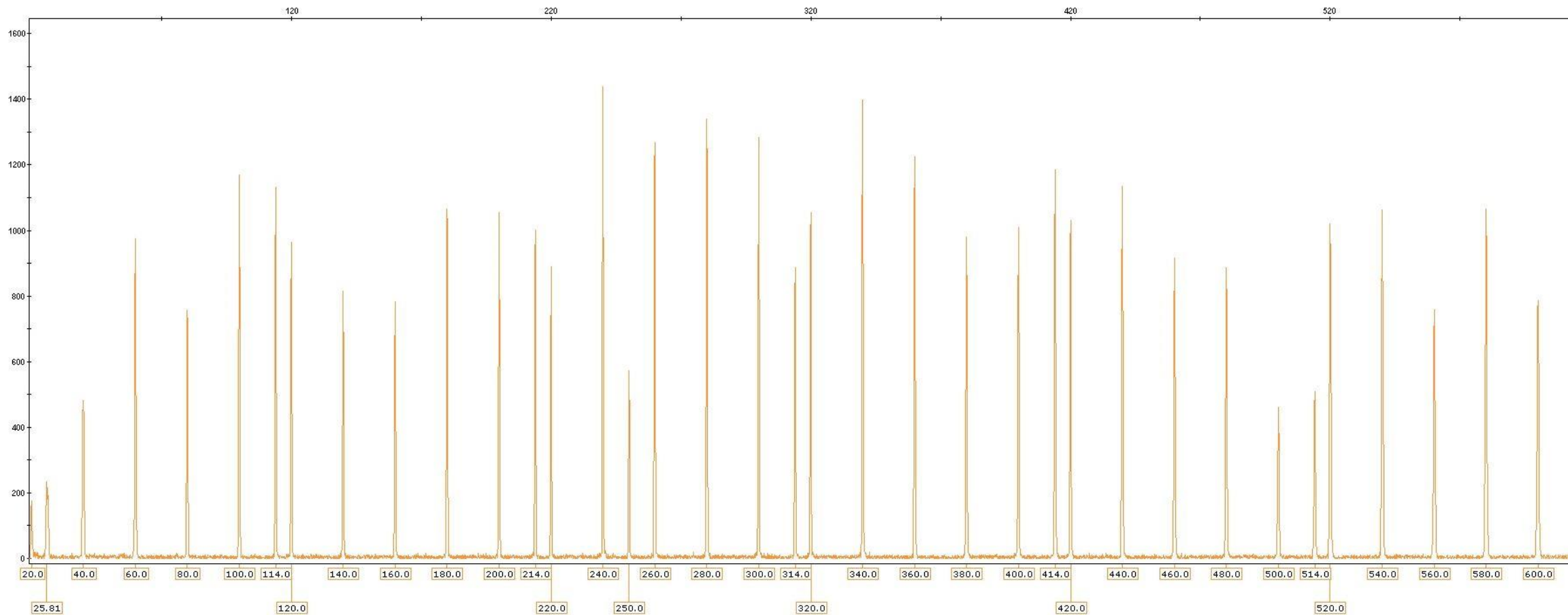
# Fragment analysis

Fluorescence capillary electrophoresis on a genetic analyzer - separation of microsatellites according to size



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# Size standard



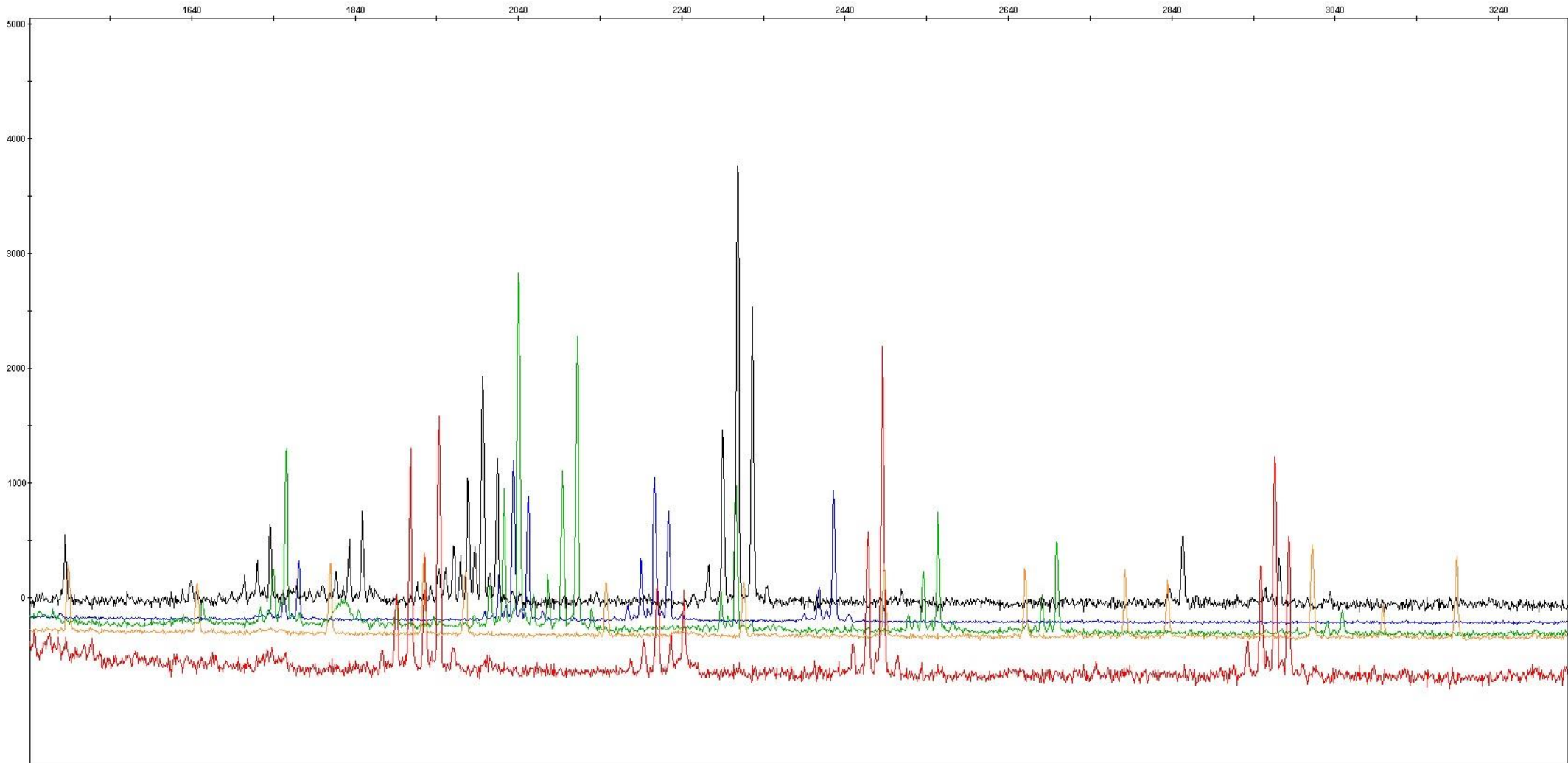
ISAGREED



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# Raw data - capillary electrophoresis

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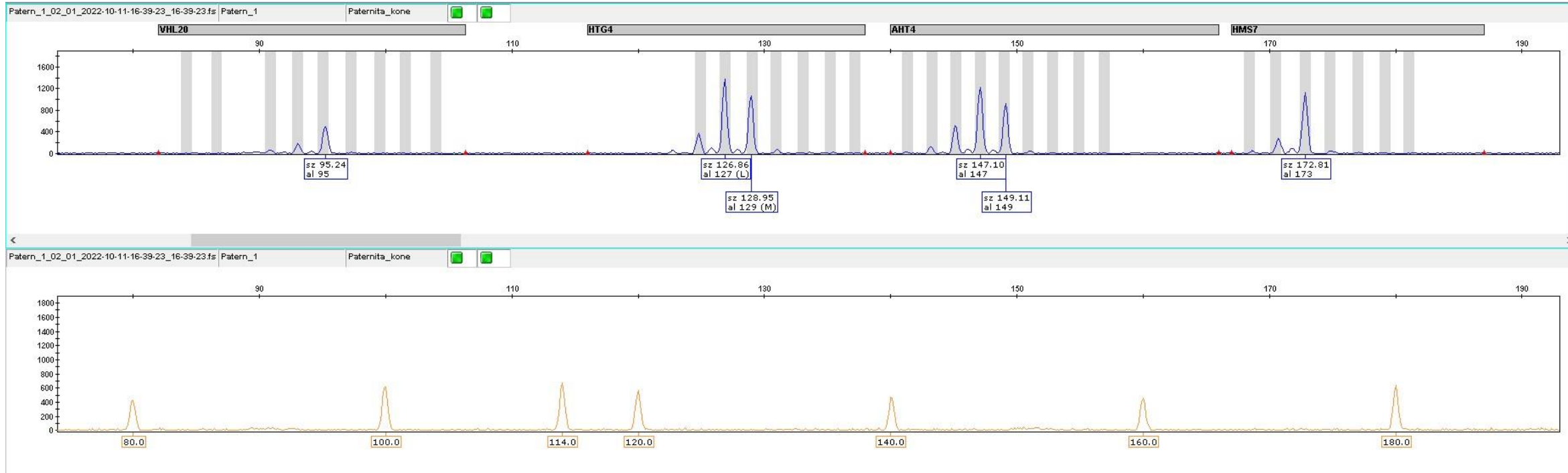


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# Evaluation - genotypes

Example of one of the colours (6-FAM) and size standard (LIZ)





# Thank you for your attention!

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