

Intra-individual difference of length heteroplamy in blood and hair shaft mitochondrial DNA

**Ukhee Chung, Hwan Young Lee, Myung Jin Park,
Ji-Eun Yoo, Gil-Ro Han, Sang-Ho Cho,
Chong-Youl Kim, Kyoung-Jin Shin**

*Department of Forensic Medicine College of Medicine, Yonsei University
Brain Korea 21 Project for Medical Science, Yonsei University
Human Identification Research Institute, Yonsei University
Department of Forensic Medicine, National Institute of Scientific Investigation*

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Mitochondrial DNA

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- General properties
 - High copy number
 - Maternal inheritance
 - Rapid rate of evolution
- D-loop; major target for forensic field
 - Hypervariable regions (HV1, HV2 and HV3)
 - Point heteroplasmy and length heteroplasmy

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Length heteroplasmy

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- Occurs in HV1, HV2, and HV3 homopolymeric C-tract
- Consequence of poor replication fidelity
- More frequent than point heteroplasmy
- Limitation to profiling of length variants

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Aims of study

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- Overcoming the limitation to analyze length heteroplasmy in mitochondrial HV2 region
- Analysis of intra-individual and inter-individual differences of length heteroplasmy
- Discussion on using length heteroplasmy for forensic identification

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Sampling and DNA extraction

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- Sampling
 - 25 individuals
 - Blood and five hair shafts
- DNA extraction
 - Using the DNA IQ™ systems (Promega) and Tissue and Hair Extraction Kit (Promega)
 - Additional decontamination step for hair shaft mtDNA extraction

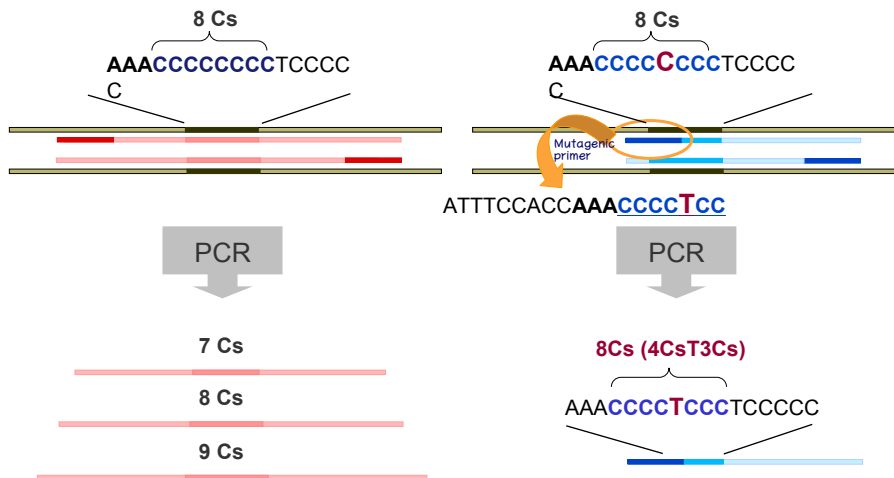
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Mutagenic primer design

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PCR amplification and detection

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- PCR amplification
 - Total volume of 10.0 ul
 - Primers
 - F291 (5'-ATTTCCACCAAACCCCTCC)
 - R389 (5'-**HEX**-CTGGTTAGGCTGGTGGTTAGG)
 - 56°C annealing temperature
 - 25 cycles for 1.0 ng of blood DNA, and 32 cycles for 0.1 ng of hair shafts mtDNA
- Size-based separation on an ABI PRISM 310 Genetic Analyzer (Applied Biosystems)

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Confirmation of mitotype

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- To examine the possibility of sample switching or contamination
- Subjects
 - Blood and hair shafts samples showing intra-individual difference in heteroplasmy mtDNA peak pattern
- Methods
 - Sequencing of the HV1 and HV2 region
 - Analysis of CA dinucleotide repeat length polymorphism in the HV3 region

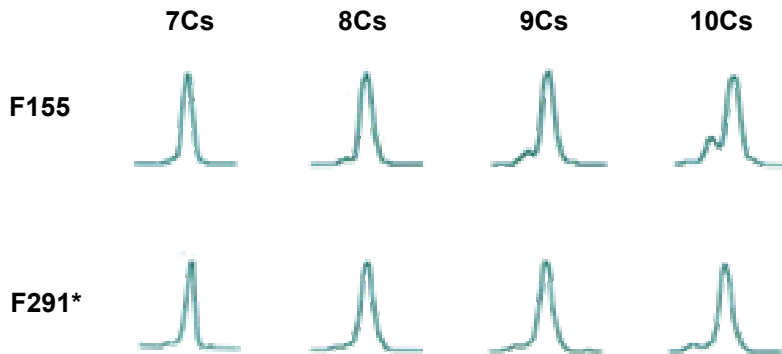
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Efficiency of mutagenic primer

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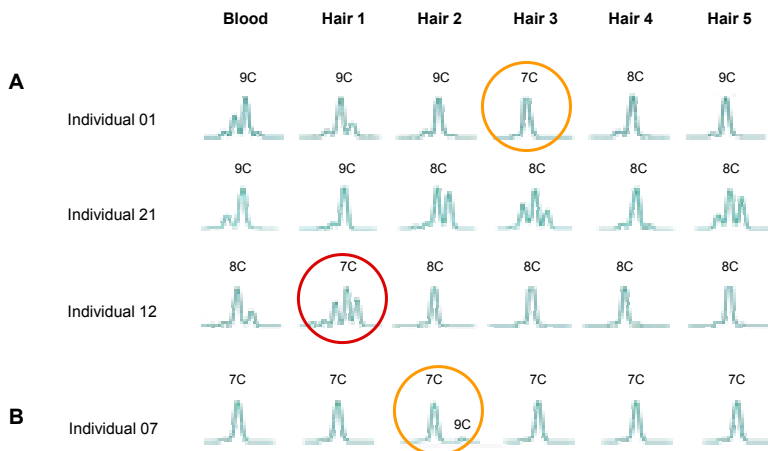
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Electropherogram of the HV2 C-tract

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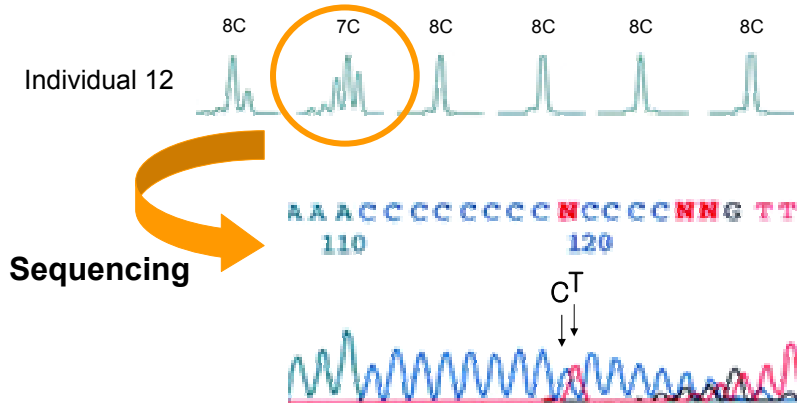
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Evidence of point heteroplasmy

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Homoplasmic and heteroplasmic individuals

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	Blood	Hair shafts		
		All homoplasmy	All heteroplasmcy	Mixed*
Homoplasmy	10	7	-	3
Heteroplasmy	15	-	13	2

* Individuals who displayed homoplasmic and heteroplasmic peak patterns in hair shafts

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Different major C-tract genotype

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Individual	Number of cytosine in major C-tract					
	Blood	Hair 1	Hair 2	Hair 3	Hair 4	Hair 5
01	9	9	9	7	8	9
10	7	7,8	8	7	7	7
12	8	7	8	8	8	8
14	9	8	9	9	8	9
16	8	8	8	9	8	8
20	8	7	8	8	7	8
21	9	9	8	8	8	8
25	9	8	9	9	9	9

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Conclusion

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Analysis of length heteroplasmy in homopolymeric C-tract of the HV2 region using primer designed to minimize stutter production

- Significant variations were observed in qualitative/quantitative peak patterns in blood and hair shaft mtDNA
- Information of length heteroplasmy in a homopolymeric C-tract of the mtDNA HV2 region cannot be used alone to support an interpretation of exclusion.

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