DUPLEX MITOCHONDRIAL DNA PCR AMPLIFICATION

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Duplex Mitochondrial DNA PCR Amplification

<u>PURPOSE:</u> To increase the amount of available mtDNA for the purposes of analysis, by performing an *in vitro* replication of template DNA using oligonucleotide primers, thermostable DNA polymerase and deoxynucleoside triphosphate bases (dNTPs) within a thermal cycler.

PROCEDURE:

A positive control, an amplification negative, and an extraction negative control (if applicable) should be included with each batch of samples being amplified to demonstrate procedural integrity. The positive control is a laboratory grade cell line, for which the mtDNA type is known.

Follow the mtDNA pre-amplification guidelines for handling the tubes and cleaning of the work surfaces. The following steps have to be performed in the appropriate dedicated areas. Evidence samples and exemplar samples should not be handled at the same time.

A. Preparing the DNA aliquots for HVI-HVII amplification

- Amplification can be performed with either Roche or Homebrew reagents.
 Homebrew reagents should be used for Missing Person's testing, body identifications, and kinship testing where the sample type is other than a hair shaft sample.
- When amplifying extracts which have nuclear DNA quantification data, the target amount of extract to be amplified is:
 - 100 pg when using Roche reagents
 - 500 pg when using Homebrew reagents
- When amplifying samples that have not been quantified (e.g., hair shaft samples), use 20ul of the extract and Roche reagents, only.
- Table I refers to the preparation of the control samples for the amplification.

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Table I – Control samples for amplification.

Sample	DNA (extract)	TE ⁻⁴
HL60 Positive Control DNA (100 pg/20 1)	20 µL	
Amplification Negative Control		20 µL
Extraction Negative Control,(s) when sample amplified neat	20 µL	
Extraction Negative Control (s) when a dilution/concentration of sample extract is amplified	Submit 20 µL of th negative at the sam dilution/concentrati concentrated than the	e extraction e on factor or more ne sample

B. Amplification Setup

- 1. For each amplification set, record the lot numbers and samples in the documentation. Label 0.2 ml PCR reaction tubes with sample label name and with date and time for the positive and negative controls.
- 2. If samples require dilution, prepare the aliquots in UV'ed 1.5mL tubes, and place the neat samples back into storage.
- 3. Master mix preparation:
 - a) For amplification using Roche reagents:
 - Prepare a Master Mix with Reaction Mix and Primer Mix, The following calculations are used:
 - Reaction Mix: number of samples N x 20 μ L Reaction Mix = μ L
 - Primer Mix: number of samples N x 10 μ L Primer Mix = μ L

Note: For ≤ 6 samples, use N, for ≥ 6 samples, use N+1. To save on reagents, individual aliquots of Reaction and Primer Mix can be made.

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- Add Reaction Mix and Primer Mix together to prepare Master Mix.
- Vortex the Master Mix and centrifuge briefly.
- $\circ \qquad \mbox{Aliquot 30 } \mu L \mbox{ of the Master Mix into the bottom of each} \\ \mbox{labeled 0.2 ml reaction tube.}$
- b) For amplification with Homebrew reagents:
 - Prepare Reaction Mix and Primer Mix master mixes according to Homebrew amplification documentation. The following amounts of reagents per sample are used:

Reaction Mix	Volume	Primer Mix	Volume
Reagents	per sample	Reagents	per sample
UltraPure dH2O	3.7 μL	UltraPure dH2O	6μL
GeneAMP 10X	5 µL	10 M HVIF	1 μL
PCR Buffer			
2.5mM dNTPs	4μL	10□M HVIR	1 μL
25mM MgCl ₂	4.8 μL	10□M HVIIF	1 μL
5U/□L AmpliTAQ Gold DNA Polymerase	2.5 µL	10□M HVIIR	1 μL
Total Reaction Mix volume per sample	20 µL	Total Primer Mix volume per sample	10 µL

The following calculations are used:

- Reaction Mix: number of samples + 1 (N + 1) x Reaction Mix Reagent Amount = $__\mu L$
- Primer Mix: number of samples + 1 (N + 1) x Primer Mix Reagent Amount = $__{\mu}L$
- Vortex the Reaction and Primer Mix and centrifuge briefly
- Aliquot 20 μ L of the Reaction Mix and 10 μ L of the primer mix into the bottom of each labeled 0.2 ml reaction tube.

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- 4. Add samples into the 0.2ml tubes. Use a sterile filter pipette tip for each sample addition. Open only one tube at a time for sample addition. The final aqueous volume in the PCR reaction mix tube will be 50 μ L. Transfer the appropriate volume of target DNA or TE⁻⁴ to each respective sample tube. After the addition of the DNA, cap each sample before proceeding to the next tube. If necessary, spin down the tubes at 1000 rcf for a few seconds.
- 5. When finished, place the rack with the 0.2ml tubes in the pre-amp room dumbwaiter. Send the samples up to the post-amp room.

C. Thermal Cycling

- 1. Turn on the Perkin Elmer 9700 Thermal Cycler.
- 2. Use the following settings to amplify the samples

9700 Thermal Cycler	The amplification file is as follows-		
Soak at 94°C for 14 minutes			
User: mtDNA	- Denature 92 °C for 15 seconds		
	34 cycles: - Anneal at 59 °C for 30 seconds		
File: lamtdna	- Extend at 72 °C for 30 seconds		
	Incubation at 72 °C for 10 minutes		
	Storage soak at 4 °C indefinitely		

- 3. Place the tubes in the tray in the heat block, slide the heated lid over the tubes, and fasten the lid by pulling the handle forward. Place the microtube rack used to set up the samples for PCR in the post-amp room bleach bath
- 4. Start the run by performing the following steps:
 - a. The main menu options are RUN CREATE EDIT UTIL USER. To select an option, press the F key directly under that menu option.
 - b. Verify that the user is set to "mtDNA" if not, select the USER option (F5) to display the "Select User Name" screen.
 - c. Use the circular arrow pad to highlight "mtDNA." Select the ACCEPT option (F1).

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d.	Press the RUN button (F1) and select the "lamtdna"	file.
e.	Verify that the reaction volume is set to 50 μ L and t 9600 (very important) .	he ramp speed is set to
f.	If all is correct, select the START option (F1).	
	The run will start when the heated cover reaches 103 °C.	The screen will then
	display a flow chart of the run conditions. A flashing lin	e indicates the step
	being performed, hold time is counted down. Cycle num	ber is indicated at the
	top of the screen, counting up.	
		1 1
g.	the appropriate instrument name.	e documentation under
h.	Upon completion of the amplification, press the STO	OP button repeatedly
	until the "End of Run" screen is displayed, and remo	ove your samples.
	Select the EXIT option (F5). Wipe any condensatio	n from the heat block
	with a Kimwipe and pull the lid closed to prevent du	ist from collecting on
	the heat block. Turn the instrument off.	
1.	After removing your samples, place them in the app	ropriate 2-8 °C
	tune (avampler, avidance, or quality control). Page	a according to sample
	when samples were amplified on the cover of the 0.2	2 mI PCR storage
	hox	2 III. I CIX Storage
RTANT:		

Turn instruments off **ONLY** when the Main Menu is displayed, otherwise there will be a Power Failure message the next time the instrument is turned on. It will prompt you to review the run history. Unless you have reason to believe that there was indeed a power failure, this is not necessary. Instead, press the STOP button repeatedly until the Main Menu appears.

In case of a real power failure the 9700 thermal cycler will automatically resume the run if the power outage did not last more than 18 hours. The Uninterruptible Power Supply (UPS) present in the amplification room will power the thermal cyclers for about 2-3 hours in the case of a total

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power outage. The history file contains the information at which stage of the cycling process the instrument stopped. Consult with the QA team and/or the Technical Leader on how to proceed.

