

FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR ANALYSIS

Amplification using the PowerPlex Y23 System		
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PowerPlex® Y23 Sample Preparation for Amplification

1 Procedure

PPY23 Sample Input Amount
Optimal - 500pg of male DNA*
Minimum – 100pg of male DNA

*The option for amplification with a greater input amount is available if determined appropriate for the sample by the analyst.

- 1.1 Retrieve the following reagents from the associated refrigerator and/or freezer and record the lot numbers.

PowerPlex® Y23 10X Primer Pair Mix
PowerPlex® Y23 5X Master Mix
Water, Amplification Grade for PPY23
2800M Control DNA for PPY23, 10ng/ul

- 1.2 Retrieve sample(s) needed for amplification from associated refrigerator and/or freezer.
- 1.3 Prepare dilutions (if necessary) in 1.5 mL tubes according to the values listed on the test batch data entry screen or the “FBAmplificationSheet”, **using Promega Amplification Grade Water for PPY23**, for each sample, according to Table 1. Vortex and centrifuge samples prior to aliquoting for dilution.

TABLE 1: Dilutions

Dilution	Amount of DNA Template (uL)	Amount of Water (uL)
0.25	3 or (2)	9 or (6)
0.2	2	8
0.1	2	18
0.05	2	38
0.04	4 or (2)	96 or (48)
0.02	2 or (1)	98 or (49)
0.01	1 or (2)	99 or (198)
0.008	4 or (2)	496 or (248)

- 1.4 Label amp tubes using the values generated by LIMS. These values can be found in the test batch output samples or on the “FBAmplificationSheet”.

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- 1.5 Centrifuge reagent tubes briefly to bring contents to the bottom and then vortex for 15 seconds before use. Do NOT re-centrifuge the Master Mix or Primer Pair Mix as this may cause the reagents to be concentrated at the bottom of the tube.
- 1.6 Consult the Reagents tab in LIMS for the exact amount of PowerPlex® Y23 10X Primer Pair Mix and PowerPlex® Y23 5X Master Mix to add.

Reagent	Per reaction
10X Primer Pair Mix	2.5 µL
5X Master Mix	5.0 µL
Mastermix total:	7.5 µL
DNA	17.5 µL

- 1.7 Vortex prepared Master Mix and all samples to be aliquoted for 5-10 seconds. After vortexing, **briefly centrifuge** master mix and samples.
- 1.8 Add **7.5 µL** of the prepared master mix to each tube that will be utilized, changing pipette tips and remixing master mix as needed.
- 1.9 **Witness Step.** Have another analyst witness the sample set-up.
 - 1.9.1 For the input samples, confirm the tube label and sample ID for each sample. For the output samples, **the entire amp tube label must be read for each sample.**
- 1.10 Positive Control for PPY23 (PPY23 2800M Control in LIMS) – total input amount of **250pg**.
 - 1.10.1 Perform dilution and aliquot positive control according to amplification sheet
- 1.11 Amplification Negative
 - 1.11.1 17.5 uL of Water, Amplification Grade for PPY23 (PPY23 H2O in LIMS)
- 1.12 Samples
 - 1.12.1 Aliquot samples according to amplification sheet
- 1.13 All amplification tubes should have a total final volume of 25uL.
- 1.14 Ensure that all caps are properly closed prior to sending the samples to the post-amplification laboratory.
- 1.15 Spin down samples at 1000 RPM for one minute.

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2 PowerPlex® Y23 Conditions for the Applied Biosystems GeneAmp PCR System 9700

- 2.1 Turn on the ABI 9700 Thermal Cycler.
- 2.2 Choose the following program in order to amplify these samples:

PowerPlex® Y23
user: casework
file: ppy23

- 2.3 PowerPlex® Y23 PCR Conditions for the Applied Biosystems GeneAmp PCR System 9700

PowerPlex® Y23 user: casework file: ppy23	The PowerPlex® Y23 file is as follows: Soak at 96°C for 2 minutes 30 Cycles : Denature at 94°C for 10 seconds : Anneal at 61°C for 60 seconds : Extend at 72°C for 30 seconds 20-minute incubation at 60°C. Storage soak indefinitely at 4°C
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- 2.4 Record instrument in LIMS
- 2.5 The run will start when the heated cover reaches temperature. The screen will then display a flow chart of the run conditions. A flashing line indicates the step being performed, hold time is counted down. Cycle number is indicated at the top of the screen, counting up.
- 2.6 Upon completion of the amplification:
 - 2.6.1 Remove samples and press the STOP button repeatedly until the “End of Run” screen is displayed.
 - 2.6.2 Select the EXIT option (F5).
 - 2.6.3 Wipe any condensation from the heat block with a lint free wipe and pull the lid closed to prevent dust from collecting on the heat block.

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- 2.6.4 Turn the instrument off.
- 2.6.5 NOTE: The 4°C storage soak step is not meant to store samples for an extended period. Samples should be removed from the instrument and placed in the 4°C refrigerator at the earliest convenience.
- 2.7 Place the microtube rack used to set-up the samples for PCR in the container of 10% bleach container in the Post-Amp area.
- 2.8 After completion of the thermal cycling protocol, store amplified product at 4°C and proceed with fragment analysis.
- 2.9 Complete the LIMS test batch:
- 2.9.1 Fill out the Performed By tab for the Test Batch Review.
- 2.9.2 Select all output samples and click Review to perform the test batch approval.
- 2.9.3 A batch reviewer will then complete the Test Batch Tech Review.
- 2.10 Schedule the samples to the appropriate STR test batch and create the test batch.

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