

INSTRUCTIONS FOR USE OF PRODUCTS DC2305 AND DC2320.

### Protocol for Amplification and Analysis of DNA from Storage Card Punches

This document is a quick protocol for experienced users to amplify DNA from storage card punches. Quick protocols are also available for amplifying extracted DNA and DNA from swabs. For complete protocol information and troubleshooting tips, see the PowerPlex® Y23 System Technical Manual #TMD035, which is available online at: www.promega.com/protocols/

### Before You Begin

The PowerPlex® Y23 System is compatible with the following sample types:

FTA® Sample Types	NonFTA Sample Types
Buccal cells collected on FTA® cards with Whatman	Buccal samples on Bode Buccal DNA Collector™ devices
EasiCollect™ or Fitzco Sampact™ devices	
Buccal cells collected with swabs transferred to	Blood and buccal samples on nonFTA card punches
FTA® or Indicating FTA® cards	(e.g., S&S 903)
Liquid blood (from collection or storage Vacutainer® tubes	
or finger sticks) snotted onto ETA® cards	

No preprocessing steps are required for FTA® card punches. For nonFTA cards, the punches must be preprocessed as described below. For complete protocol information see the *PunchSolution™ Kit Technical Manual #*TMD038, which is available online at: www.promega.com/protocols/

- Add 10µl of PunchSolution™ Reagent (Cat.# DC9271) to one 1.2mm punch placed in a well of the 96-well reaction plate.
   Note: Do not cover the plate or place the plate in a thermal cycler with a closed, heated lid.
- 2. Incubate plate at 70°C for 30 minutes or until wells are dry.

#### **PCR Setup**

- 1. Thaw all pre-amplification components just prior to use.
- 2. Centrifuge tubes briefly, then vortex for 15 seconds before each use. Do not centrifuge after vortexing.
- 3. Determine the number of reactions including positive and negative controls. Add 1 or 2 reactions to this number.
- 4. Prepare the PCR amplification mix by combining the components as shown below.

Component	volume per Reaction	×	Number of Reactions	=	Volume
Water, Amplification Grade	17.5µl	×		=	
PowerPlex® Y23 5X Master Mix	5.0µl	×		=	
PowerPlex® Y23 10X Primer Pair Mix	2.5μΙ	×		=	
Total volume	25μΙ				

- 5. Vortex the PCR amplification mix for 5-10 seconds.
- 6. For FTA® storage cards, add one or two 1.2mm punches from a card containing a buccal sample or one 1.2mm punch from a card containing whole blood to the appropriate wells of the reaction plate. For nonFTA card punches, add the PCR amplification mix to the PunchSolution™ Reagent-treated punches. It also is acceptable to add the FTA® card punch first, then add the PCR amplification mix.



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- For the positive amplification control, vortex the tube of 2800M Control DNA, then dilute an aliquot to 5.0ng/µl and add 1µl to a
  reaction well containing 25µl of PCR amplification mix.
- 8. Reserve a well containing PCR amplification mix as a negative amplification control.
- 9. Seal the plate, and briefly centrifuge the plate.

### Thermal Cycling

The PowerPlex® Y23 System is designed for use with the GeneAmp® PCR System 9700 with a silver or gold-plated silver sample block with Max mode as the ramp speed.

 Program the thermal cycler with the following conditions. Refer to the technical manual for more information. For DNA from punches, we recommend using 26 cycles. Optimize the cycle number as required.

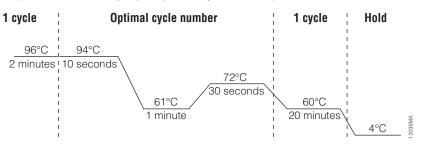


Figure 1. The thermal cycling protocol for the GeneAmp® PCR System 9700 thermal cycler.

**Optional:** Record the cycle number as optimized in your laboratory.

2. Proceed with the analysis, or store amplified samples at -20°C in a light-protected box until ready to analyze.

#### Additional Notes:



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### Instrument Setup and Sample Preparation



A passing spectral calibration must be generated using the PowerPlex® 5C Matrix Standard (Cat.# DG4850) prior to sample analysis. See the *PowerPlex® 5C Matrix Standard Technical Manual* #TMD049 for more information.

#### Instrument Setup

- For the Applied Biosystems® 3500 or 3500xL Genetic Analyzer, we recommend preheating the oven at 60°C for at least 30 minutes prior to the first injection.
- 2. Use the following parameters when setting up the instrument. Refer to the instrument user's manual for additional details.

			Injection	
Genetic Analyzer	Run Module	Dye Set	Parameters <sup>1</sup>	Run Time
Applied Biosystems® 3500	HID36_P0P4	Promega G5	1.2kV, 15 seconds	1,210-1,500 seconds
Applied Biosystems® 3500xL	HID36_P0P4	Promega G5	1.2kV, 24 seconds	1,210-1,500 seconds
Applied Biosystems® 3130 and 3130x/	HIDFragmentAnalysis36_POP4	G5 <sup>2</sup>	3kV, 5 seconds	1,500 seconds
ABI PRISM® 3100 and 3100-Avant	HIDFragmentAnalysis36_POP4	G5 <sup>2</sup>	3kV, 5 seconds	1,500 seconds

Injection time may be modified (2–24 seconds) to increase or decrease the observed peak heights.

**Optional:** Record the injection conditions as optimized in your laboratory.

#### Additional Notes:

<sup>&</sup>lt;sup>2</sup>Confirm that the active dye set is the file generated for the PowerPlex® 5-dye chemistry.



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### **Instrument Setup and Sample Preparation (continued)**

### Sample Preparation

Prepare samples for capillary electrophoresis immediately before loading.

- Centrifuge the WEN Internal Lane Standard 500 Y23 (WEN ILS 500 Y23) briefly, then vortex for 15 seconds before each use. Do not centrifuge after vortexing.
- 2. Calculate the number of samples including the number of allelic ladders per run. Add 1 or 2 reactions to this number.
- Prepare a loading cocktail by combining and mixing the WEN ILS 500 Y23 and Hi-Di™ formamide. You may need to optimize the
  volume of WEN ILS 500 Y23.

Component	Volume per Sample	Number of × Samples	=	Final Volume
WEN ILS 500 Y23	0.5µl¹ >	×	=	
Hi-Di™ formamide	9.5µI >	×	=	

<sup>&</sup>lt;sup>1</sup>The volume of WEN ILS 500 Y23 can be adjusted to change the intensity of the size standard peaks based on laboratory preferences.

Optional: Record the volume of WEN ILS 500 Y23 per sample as optimized in your laboratory.

- 4. Vortex the loading cocktail for 10-15 seconds, and pipet 10µl of formamide/internal lane standard mix into each well.
- Add 1µI of amplified sample (or 1µI of PowerPlex® Y23 Allelic Ladder Mix). Cover wells with appropriate septa, and centrifuge plate briefly.
- Denature samples at 95°C for 3 minutes, then immediately chill on crushed ice or in an ice-water bath for 3 minutes. Denature samples just prior to loading the instrument.
- 7. Place the plate assembly on the autosampler.
- 8. Start the capillary electrophoresis run.

### **Data Analysis**

The panels, bins and stutter text files needed for data analysis using GeneMapper® ID software, version 3.2, and GeneMapper® ID-X software, version 1.2 or higher, are available for download at:

www.promega.com/resources/software-firmware/genemapper-id-software-panels-and-bin-sets/

For complete protocol information see the *PowerPlex® Y23 System Technical Manual #*TMD035, available online at: www.promega.com/protocols/

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