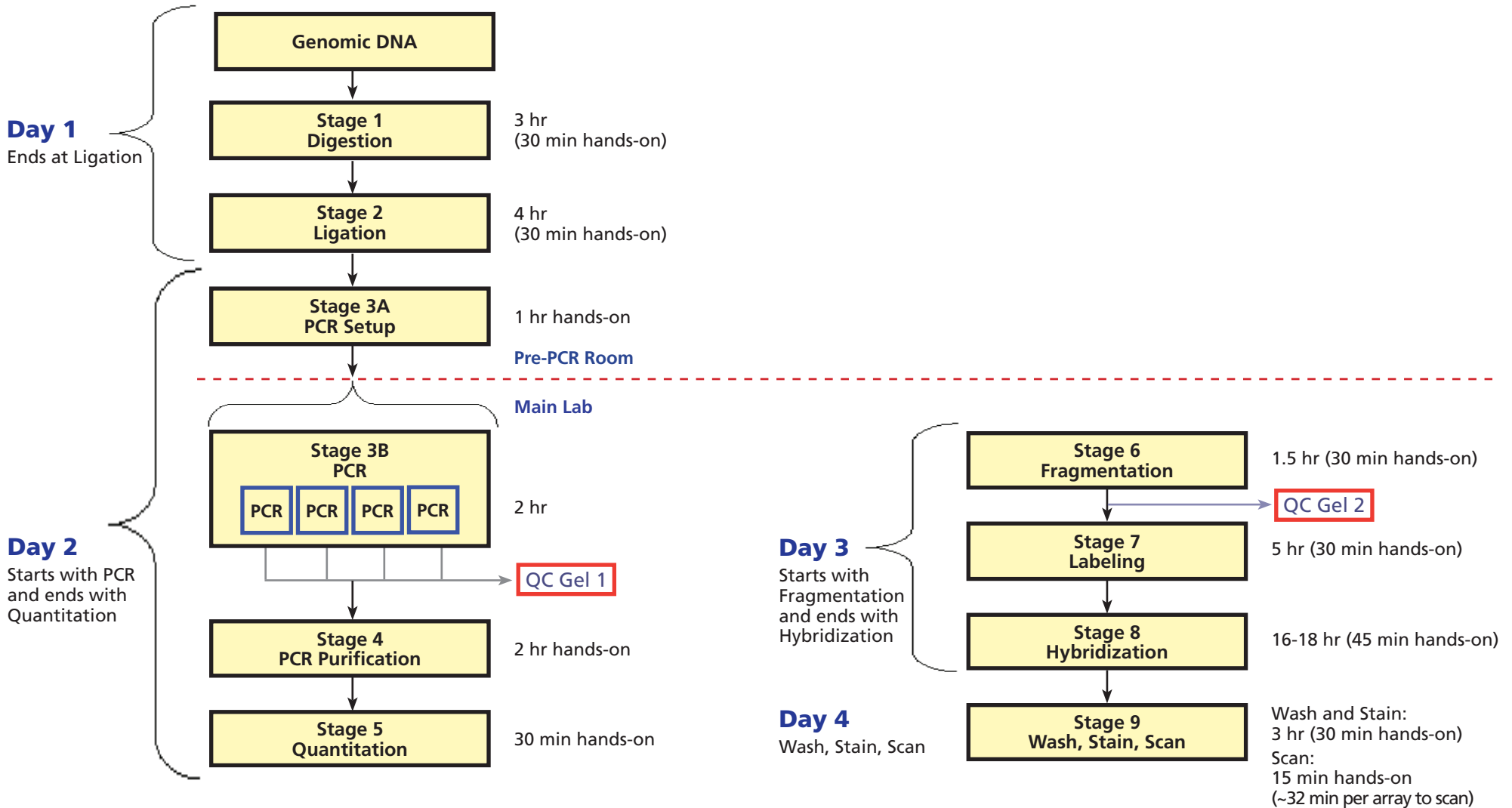


# Quick Reference Card

## CytoScan™ Assay 24 Samples: Workflow Overview

The Affymetrix® CytoScan™ Assay protocol is optimized for processing 8 to 24 samples at a time to obtain whole genome copy number and SNP information from Affymetrix® CytoScan™ Arrays. The CytoScan™ Assay protocol supports processing of as little as eight samples, two of which are a positive and negative control. This protocol is intended for research use only and is not intended for diagnostic purposes.



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# Quick Reference Card

## CytoScan™ Assay 24 Samples: Stage 1 – Digestion

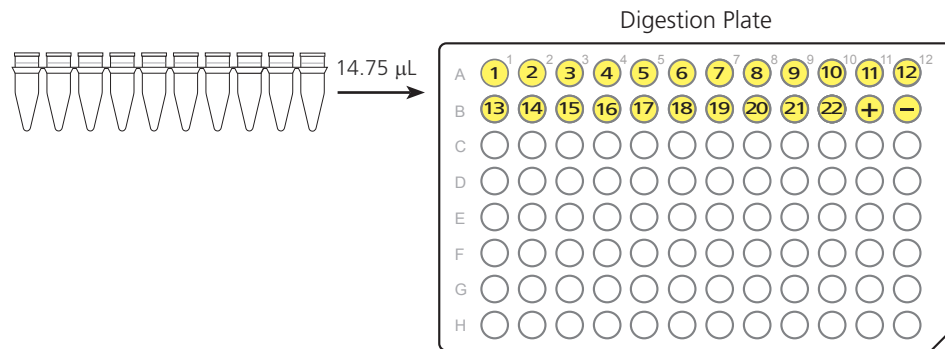
1. Add gDNA to wells marked 1 through 22 in the plate diagram.
2. Thaw 10X Nsp I Buffer and 100X BSA at room temperature. Vortex and spin down, then place on ice.
3. Leave Nsp I enzyme at -20 °C until ready to use.
4. Add 5 µL of the Genomic DNA, supplied in the kit as positive control, to the well marked "+".
5. Add 5 µL of Low EDTA TE as negative control to the well marked "-".
6. Prepare the Digestion Master Mix.

10. Ensure that the lid of the thermal cycler is preheated.
11. Seal the plate with an adhesive film.
12. Vortex the plate at high speed in 5 sector format, 1 sec per sector.
13. Spin down at 2000 rpm for 1 min.
14. Load the plate onto the thermal cycler and run the *CytoScan Digest* program.

Temp	Time
37 °C	2 hr
65 °C	20 min
4 °C	Hold

Reagent	Per Sample	24 Samples MM (with 20% overage)	✓	Lot Number
Chilled Affymetrix® Nuclease-Free Water	11.55 µL	332.6 µL		
10X Nsp I Buffer	2.00 µL	57.6 µL		
100X BSA	0.20 µL	5.8 µL		
Nsp I	1.00 µL	28.8 µL		
<b>Total Volume</b>	<b>14.75 µL</b>	<b>424.8 µL</b>	—	—

7. Vortex the Digestion Master Mix at high speed 3 times, 1 sec each time, and spin down.
8. Aliquot the Digestion Master Mix equally to strip tubes.
9. Use a multi-channel pipette to add 14.75 µL to the samples.



Samples	Volume/Sample
gDNA (50 ng/µL)	5.00 µL (250 ng)
Digestion Master Mix	14.75 µL
<b>Total Volume</b>	<b>19.75 µL</b>

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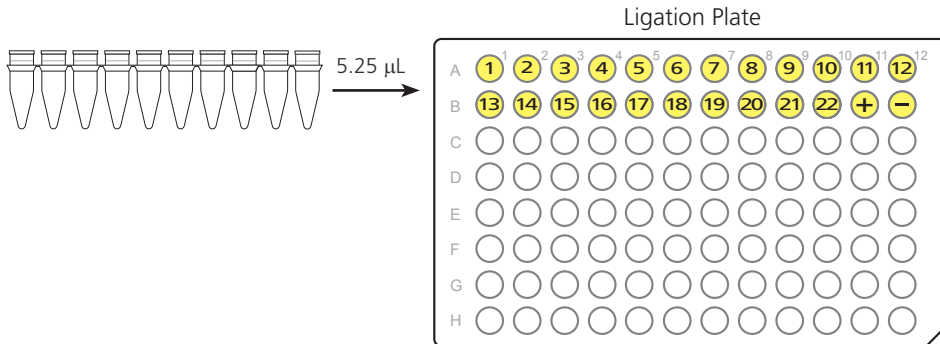
# Quick Reference Card

## CytoScan™ Assay 24 Samples: Stage 2 – Ligation

1. Thaw the 10X T4 DNA Ligase Buffer and 50 µM Adaptor, Nsp I at room temperature. Vortex to ensure any precipitate is resuspended and the buffer is clear. Place on ice.
2. Leave T4 DNA Ligase at -20 °C until ready to use.
3. Prepare the Ligation Master Mix

Reagent	Per Sample	24 Samples MM (with 25% overage)	✓	Lot Number
10X T4 DNA Ligase Buffer	2.50 µL	75.0 µL		
50 µM Adaptor, Nsp I	0.75 µL	22.5 µL		
T4 DNA Ligase	2.00 µL	60.0 µL		
<b>Total Volume</b>	<b>5.25 µL</b>	<b>157.5 µL</b>	—	—

4. Vortex the Ligation Master Mix at high speed 3 times, 1 sec each time, and spin down.
5. Aliquot the Ligation Master Mix equally to strip tubes.
6. Use a multi-channel pipette to add 5.25 µL to the samples.



Samples	Volume/Sample
Digested Sample	19.75 µL
Ligation Master Mix	5.25 µL
<b>Total Volume</b>	<b>25.00 µL</b>

7. Seal the plate with an adhesive film.
8. Vortex the plate at high speed in 5 sector format, 1 sec per sector.
9. Spin down at 2000 rpm for 1 min.
10. Load the plate onto the thermal cycler and run the *CytoScan Ligase* program.

Temp	Time
16 °C	3 hr
70 °C	20 min
4 °C	Hold

11. Proceed to the PCR setup.

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# Quick Reference Card

## CytoScan™ Assay 24 Samples: Stage 3A – PCR

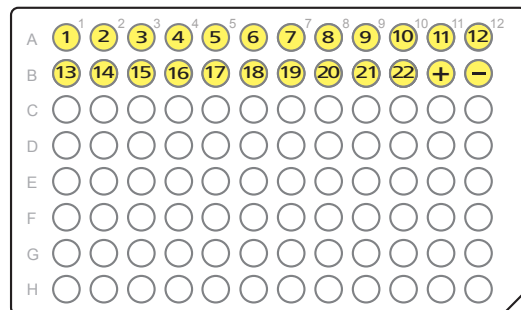
1. Ensure that the ligation plate is sealed properly.
2. Spin down at 2000 rpm for 1 min.
3. Dilute the ligated samples.

Samples	Volume/Sample
Ligated Samples	25 $\mu$ L
Chilled Affymetrix® Nuclease-Free Water	75 $\mu$ L
<b>Total Volume</b>	<b>100 <math>\mu</math>L</b>



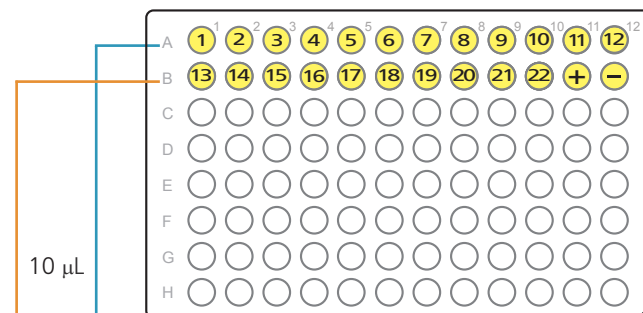
Chilled Affymetrix®  
Nuclease-Free Water

75  $\mu$ L

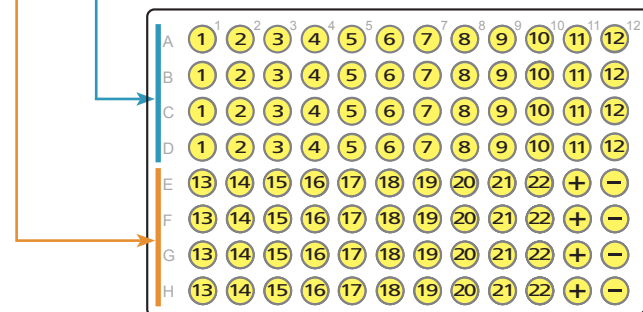


Diluted Ligated Samples

4. Seal the plate with an adhesive film.
5. Vortex at high speed in 5 sector format, 1 sec per sector. **REPEAT** vortexing, then spin down at 2000 rpm for 1 min.
6. Transfer four 10  $\mu$ L aliquots of each sample to the PCR plate.



Diluted Ligated Samples



PCR Plate

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# Quick Reference Card

## CytoScan™ Assay 24 Samples: Stage 3A – PCR (continued)

7. Thaw the 10X TITANIUM™ Taq PCR Buffer, dNTP Mixture, PCR Primer 002, and diluted ligated samples at room temperature. After thawing, immediately place on ice. Vortex and spin down all reagents.
8. Keep the GC-Melt Reagent and Affymetrix® Nuclease-Free water on ice.
9. Leave the 50X TITANIUM™ Taq DNA Polymerase at -20 °C until ready to use.
10. Prepare the PCR Master Mix in a 50 mL centrifuge tube. Assemble the master mix on ice.

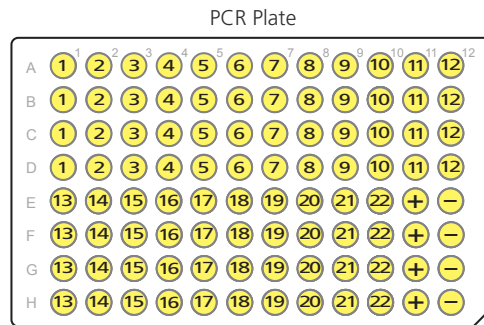
Reagent	Per Sample	24 Samples MM (with 15% overage)	✓	Lot Number
Chilled Affymetrix® Nuclease-Free Water	39.5 µL	4360.8 µL		
10X TITANIUM™ Taq PCR Buffer	10.0 µL	1104.0 µL		
GC-Melt Reagent	20.0 µL	2208.0 µL		
dNTP Mixture (2.5 mM each)	14.0 µL	1545.6 µL		
PCR Primer 002	4.5 µL	496.8 µL		
50X TITANIUM™ Taq DNA Polymerase	2.0 µL	220.8 µL		
<b>Total Volume</b>	<b>90.0 µL</b>	<b>9936.0 µL</b>	—	—

11. Vortex the PCR Master Mix at high speed 3 times, 1 sec each.
12. Transfer the PCR Master Mix to a reservoir.
13. Use a multi-channel pipette to add 90 µL to the samples.



PCR Master Mix

90 µL →



Sample	Volume/Sample
Ligated Sample	10 µL
PCR Master Mix	90 µL
<b>Total Volume</b>	<b>100 µL</b>

14. Seal the PCR plate, vortex at high speed in 5 sector format, 1 sec per sector. REPEAT vortexing, then spin down at 2000 rpm for 1 min.
15. Keep the plate on ice and move to the Post-PCR Room/Area.

### Pre-PCR Area

### Post-PCR Area

16. Ensure that the thermal cycler lid is preheated.
17. Load the plate onto the thermal cycler and run the *CytoScan PCR* program.
18. Hold at 4 °C.

### CytoScan PCR – ABI 9700

Temp	Time	Cycles
94 °C	3 min	—
94 °C	30 sec	30
60 °C	45 sec	
68 °C	15 sec	
68 °C	7 min	—
4 °C	Hold	—

Volume: 100 uL  
Specify *Maximum* mode

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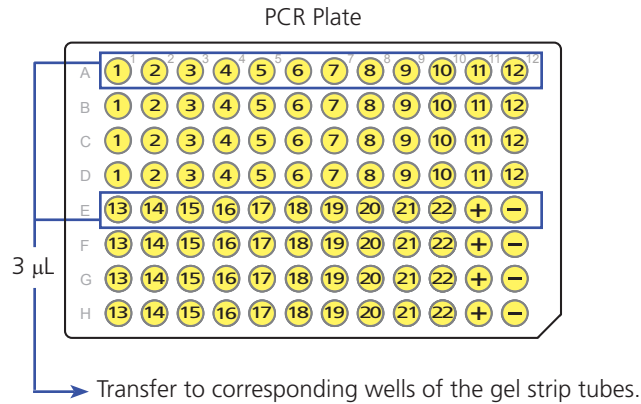
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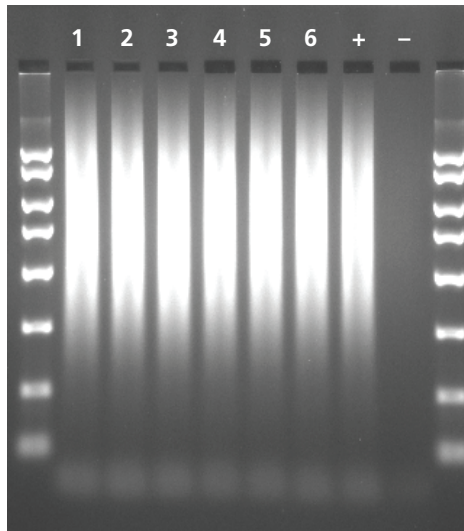
# Quick Reference Card

## CytoScan™ Assay 24 Samples: Stage 3B – PCR Product Check

1. Aliquot 5  $\mu$ L of Affymetrix® Nuclease-Free water and 2  $\mu$ L of USB 5X RapidRun™ Loading Dye into 24 wells of new gel strip tubes.
2. Transfer 3  $\mu$ L of each PCR product from one row to the corresponding wells of the gel strip tubes.
3. Seal the gel strip tubes.



4. Vortex and spin down.
5. Load 8  $\mu$ L of the sample mix from the gel strip tubes onto a 2% agarose gel. Load 5  $\mu$ L of USB 50-2000 bp Ladder in the lanes before and after the samples. Run the gel at 5V/cm for 45 min.



Example of PCR products run on a 2% TBE gel at 5 V/cm for 45 min.  
Majority of product should be between 150 and 2000 bp.

6. While the gel is running, begin Stage 4 – PCR Purification.

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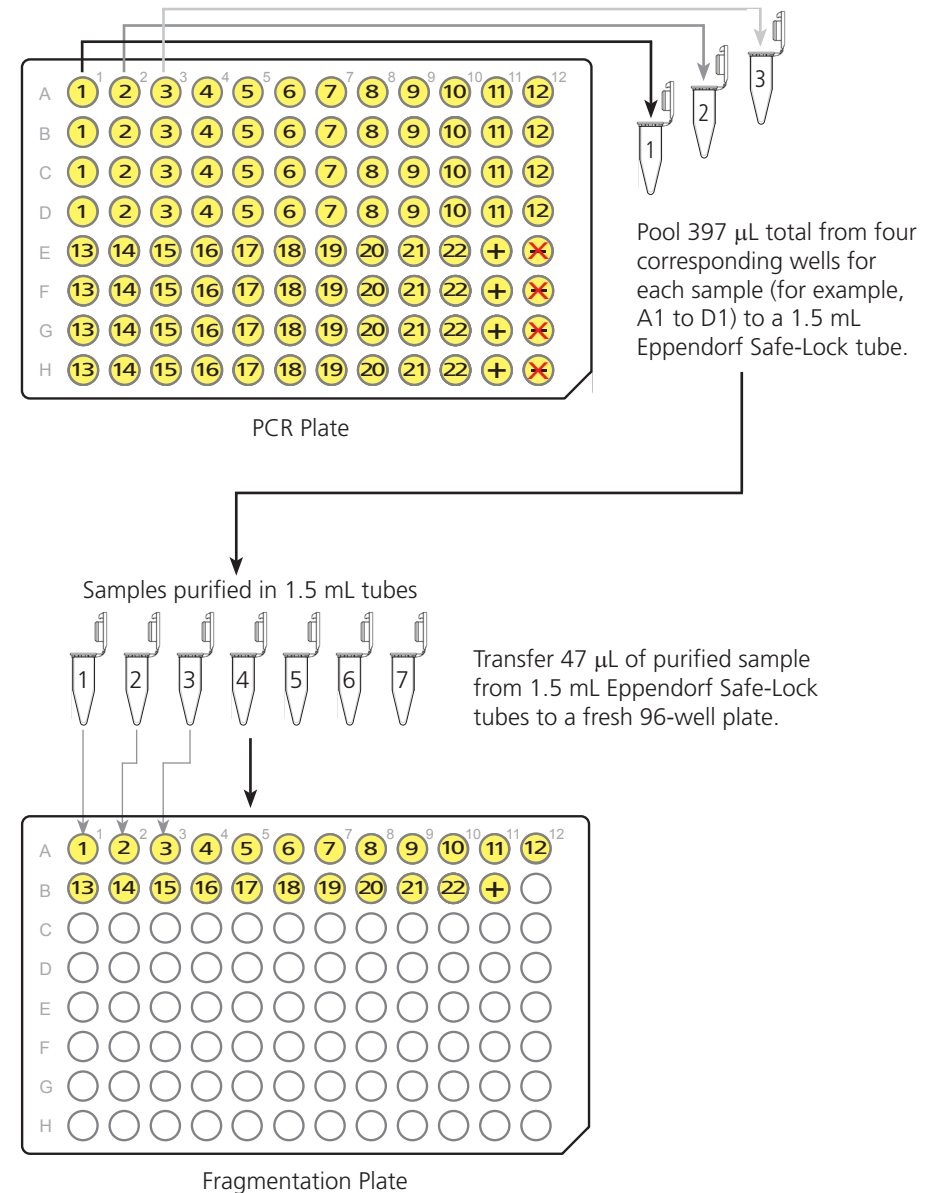
# Quick Reference Card

## CytoScan™ Assay 24 Samples: Stage 4 – PCR Product Purification

- Pool all 4 PCR products for each sample by transferring all PCR reactions to the appropriately marked 1.5 mL Eppendorf Safe-Lock tube.
- Examine the PCR plate to ensure all of the volume from each well has been transferred.
- Thoroughly mix the Purification Beads by inverting the bottle up and down until the mixture is homogeneous.
- Add 720  $\mu$ L of Purification Beads to each pooled sample. If using a multi-channel P1000 pipette, first add 21 mL of Purification Beads to a reservoir, then take 720  $\mu$ L of beads from the reservoir and add to the samples.
- Securely cap each tube and mix well by inverting 10 times.
- Incubate at room temperature for 10 min.
- Centrifuge the tubes—with hinges facing out—for 3 min at maximum speed (16,100 rcf).
- Place the tubes on a magnetic stand (for example, MagnaRack™).
- Leaving the tubes in the stand, pipet off the supernatant without disturbing the bead pellet. Discard the supernatant.

**NOTE:** Be sure to add 45 mL of absolute ethanol to the Purification Wash Buffer prior to use.

- Using a P1000 pipet, add 1.0 mL Purification Wash Buffer to each tube.
- Cap the tubes, load into the foam adapter, and vortex at maximum setting for 2 min.
- Centrifuge the tubes for 3 min at 16,100 rcf with hinges facing out.
- Place the tubes back on the magnetic stand.
- Leaving tubes in the stand, pipet off the supernatant without disturbing the bead pellet. Discard the supernatant.
- Spin the tubes for 30 sec at 16,100 rcf with hinges facing out, then place them back on the magnetic stand.
- Using a P20 pipet, remove any drops of Purification Wash Buffer from the bottom of each tube.
- Allow any remaining Purification Wash Buffer to evaporate by taking the tubes OFF the magnetic stand and leaving them UNCAPPED at room temperature for 10 min.
- Using a P200 pipet, add 52  $\mu$ L of Elution Buffer to each tube, directly dispensing onto the beads.
- Cap the tubes, load into the foam adapter, and vortex at maximum power for 10 min to resuspend the beads.
- If the beads are not fully resuspended, flick the tubes to dislodge the pellet and vortex an additional 2 min.
- Centrifuge the tubes for 3 min at 16,100 rcf with hinges facing out.
- Place the tubes on the magnetic stand for 10 min until all beads are pulled to the side.
- Transfer 47  $\mu$ L of eluted sample to the appropriate well of a fresh 96-well plate.
- Seal the plate tightly. Vortex at high speed for 1 sec each in all corners and in the center. Spin down at 2000 rpm for 1 min.



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## CytoScan™ Assay 24 Samples: Stage 5 – Quantitation

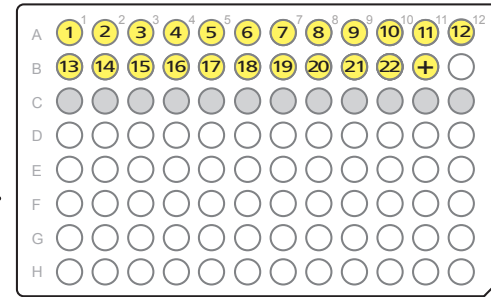
### Prepare the Quantitation Plate

1. Aliquot 198 µL of Affymetrix® Nuclease-Free water into a UV plate.
2. Add 2 µL of each purified sample.
3. Seal the plate, vortex, and spin down.

### Plate Spectrophotometer

1. Measure the OD of each PCR product at 260, 280 and 320 nm.
2. Determine the OD<sub>260</sub> measurement for the water blank and average. ....>
3. Calculate one OD reading for every sample:  
 $OD = (\text{sample OD}) - (\text{average water blank OD})$
4. Calculate the undiluted concentration for each sample in µg/µL:  $OD \times 0.05 \mu\text{g}/\mu\text{L} \times 100$ .

UV Transparent Plate

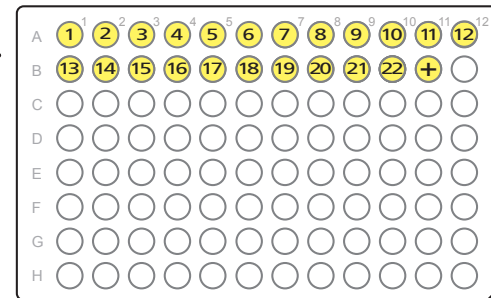


- = 198 µL Affymetrix® Nuclease-Free water + 2 µL sample
- = 200 µL Affymetrix® Nuclease-Free water for blank

### Nanodrop

1. Aliquot 18 µL of Affymetrix® Nuclease-Free water to the corresponding wells of a 96-well plate.
2. Using a P20 pipette, transfer 2 µL of each purified sample to the corresponding well of a 96-well plate. ....>
3. Seal the plate, vortex, and spin down.
4. Blank the NanoDrop using Affymetrix® Nuclease-Free water.
5. Take 2 µL of diluted sample and measure the OD of each PCR product at 260, 280 and 320 nm.
6. Calculate the undiluted concentration of each sample in µg/µL:  $(\text{concentration in ng}/\mu\text{L} \times 10) \div 1000$ .

96-well PCR Plate



- = 18 µL Affymetrix® Nuclease-Free water + 2 µL sample

### Assess OD Readings

- The average purification yield for 7 or more samples should be  $\geq 3.0 \mu\text{g}/\mu\text{L}$ . We do not recommend further processing of samples with yields  $< 2.5 \mu\text{g}/\mu\text{L}$ . If the yields are outside of the range indicated here, please consult the troubleshooting section of the *CytoScan™ Assay User Manual* (P/N 703038).
- The OD<sub>260</sub>/OD<sub>280</sub> ratio should be between 1.8 and 2.0.
- The OD<sub>320</sub> measurement should be very close to zero ( $< 0.1$ ).

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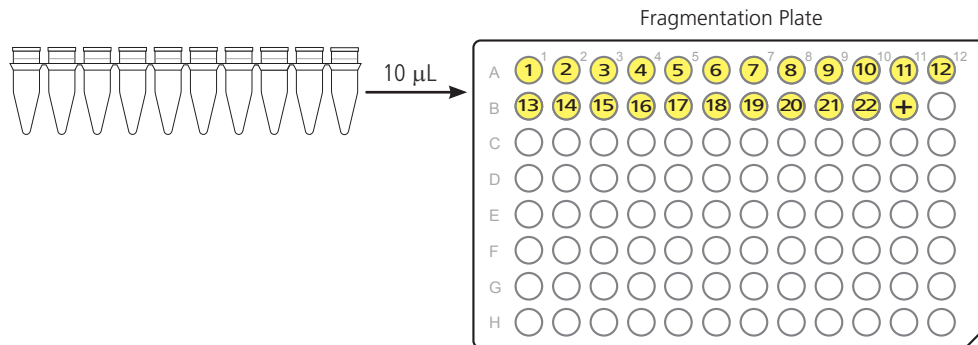
# Quick Reference Card

## CytoScan™ Assay 24 Samples: Stage 6A – Fragmentation

1. Turn down the plate centrifuge to 4 °C before proceeding into the fragmentation step.
2. Turn on the thermal cycler to pre-heat the lid.
3. Remove the plate of purified, quantitated samples from –20 °C storage and thaw. Seal the plate tightly, then vortex and spin down. Place the plate on the lower half of the cooling block on ice and chill for 10 min prior to use.
4. Leave the Fragmentation Reagent at –20 °C until ready to use.
5. Keep all reagents, including water, on ice. Perform all additions on ice.
6. Prepare the Fragmentation Master Mix as per the concentration on the enzyme tube label.

Reagent	Fragmentation Reagent Concentration					✓	Lot Number
	2.0 U/μL	2.25 U/μL	2.5 U/μL	2.75 U/μL	3.0 U/μL		
Chilled Affymetrix® Nuclease-Free Water	122.4 μL	123.2 μL	123.8 μL	124.4 μL	124.8 μL		
10X Fragmentation Buffer	158.4 μL	158.4 μL	158.4 μL	158.4 μL	158.4 μL		
Fragmentation Reagent	7.2 μL	6.4 μL	5.8 μL	5.2 μL	4.8 μL		
<b>Total Volume</b>	<b>288.0 μL</b>	<b>288.0 μL</b>	<b>288.0 μL</b>	<b>288.0 μL</b>	<b>288.0 μL</b>	—	—

7. Vortex the master mix at high speed 3 times, 1 sec each time.
8. Aliquot the Fragmentation Master Mix equally to strip tubes.
9. Using a multi-channel pipette, add 10 μL of Fragmentation Master Mix to each sample.



Samples	Volume/Sample
Purified PCR Product	45 μL
Fragmentation Master Mix	10 μL
<b>Total Volume</b>	<b>55 μL</b>

10. Seal the sample plate with an adhesive film.
11. Vortex at high speed in 5 sector format, 1 sec per sector.
12. Spin down at 2000 rpm for 1 min in a **pre-chilled centrifuge**.
13. Ensure that the thermal cycler block is preheated.
14. Load the plate onto the thermal cycler and run the *CytoScan Fragment* program. . . . . →
15. Proceed immediately to Fragmentation QC Gel.

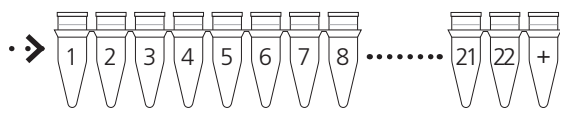
Temp	Time
37 °C	35 min
95 °C	15 min
4 °C	Hold

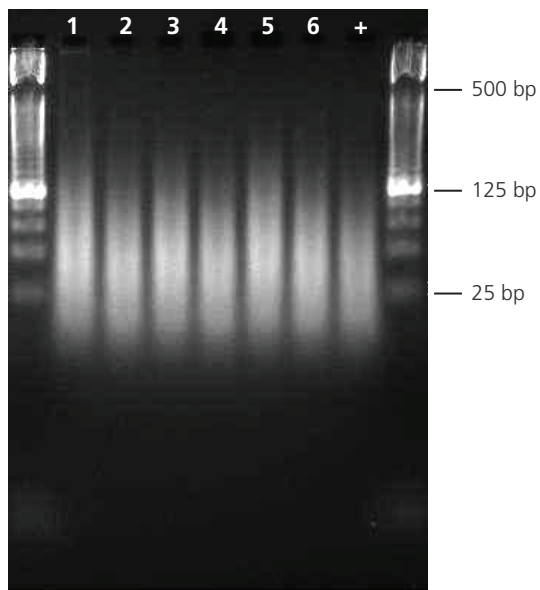
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## CytoScan™ Assay 24 Samples: Stage 6B – Fragmentation QC Gel

1. Transfer 4  $\mu\text{L}$  of each fragmented sample into strip tubes and label as Fragmentation QC Samples. . . . .  $4 \mu\text{L}$  
2. Add 28  $\mu\text{L}$  of Affymetrix® Nuclease-Free water to each strip tube. Seal the strip, vortex, and spin down.
3. Take an 8  $\mu\text{L}$  aliquot out and add to a strip tube, labeled as Gel Analysis. Add 2  $\mu\text{L}$  of USB 5X RapidRun™ Loading Dye to each tube of the strip. Seal the strip tubes tightly, vortex, and spin down.
4. Load 8  $\mu\text{L}$  of the samples onto a 4% TBE gel. Load 2  $\mu\text{L}$  of TrackIt™ 25 bp DNA Ladder before and after the samples to the first and last lanes.



Example of fragmented samples run on a 4% TBE gel at 5 V/cm for 45 min.  
Average fragment distribution is between 25 to 125 bp.

5. Run the gel at 5V/cm for 45 min.
6. Inspect the gel and compare against the figure shown above.
7. Store the remaining 24  $\mu\text{L}$  aliquot of the Fragmentation QC samples at  $-20 \text{ }^\circ\text{C}$  for further analysis on the Bioanalyzer.
8. If the QC results are good, proceed to Stage 8 – Labeling.

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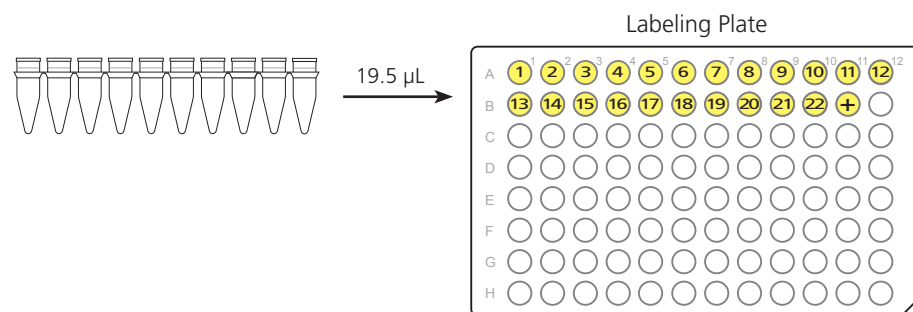
# Quick Reference Card

## CytoScan™ Assay 24 Samples: Stage 7 – Labeling

1. Thaw the 5X TdT Buffer and 30 mM DNA Labeling Reagent at room temperature, then place on ice.
2. Leave the TdT enzyme at –20 °C until ready to use.
3. Prepare the Labeling Master Mix.

Reagent	Per Sample	24 Samples MM (with 20% overage)	✓	Lot Number
5X TdT Buffer	14.0 µL	403.2 µL		
30 mM DNA Labeling Reagent	2.0 µL	57.6 µL		
TdT	3.5 µL	100.8 µL		
<b>Total Volume</b>	<b>19.5 µL</b>	<b>561.6 µL</b>	—	—

4. Vortex the Labeling Master Mix and spin down.
5. Aliquot the Labeling Master Mix equally to strip tubes.
6. Use a multi-channel pipette to add 19.5 µL to samples.



Samples	Volume/Sample
Fragmented DNA (less 4.0 µL for gel analysis)	51 µL
Labeling Mix	19.5 µL
<b>Total Volume</b>	<b>70.5 µL</b>

7. Tightly seal the plate, and vortex in 5 sector format, 1 sec per sector.
8. Spin down at 2000 rpm for 1 min.
9. Load the plate onto the thermal cycler and run the *CytoScan Label* program.

Temp	Time
37 °C	4 hr
95 °C	15 min
4 °C	Hold

10. If not proceeding to hybridization, store the Labeling plate overnight at –20 °C. Otherwise, you can also hold the Labeling Plate at 4 °C overnight.

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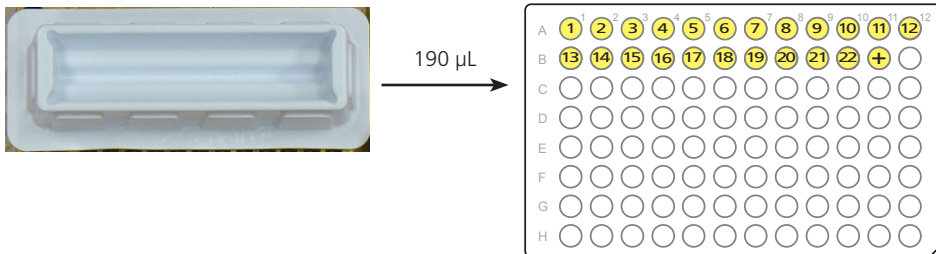
## CytoScan™ Assay 24 Samples: Stage 8 – Hybridization

1. Unpack the arrays and allow to equilibrate to room temperature prior to use.
2. Preheat the hybridization ovens for at least 1 hr at 50 °C with the rotation turned on.
3. Create a Batch Registration file.
4. Prepare the Hybridization Master Mix in a 15 mL conical tube on ice.

Reagent	Per Sample	24 Samples MM (with 20% overage)	✓	Lot Number
Hyb Buffer Part 1	165.0 µL	4752.0 µL		
Hyb Buffer Part 2	15.0 µL	432.0 µL		
Hyb Buffer Part 3	7.0 µL	201.6 µL		
Hyb Buffer Part 4	1.0 µL	28.8 µL		
Oligo Control Reagent 0100	2.0 µL	57.6 µL		
<b>Total Volume</b>	<b>190.0 µL</b>	<b>5472.0 µL</b>	—	—

5. Mix well by vortexing the master mix at high speed 3 times, 3 seconds each and pour it into a reservoir on the cold block.
6. Add 190 µL of Hybridization Master Mix to each sample.

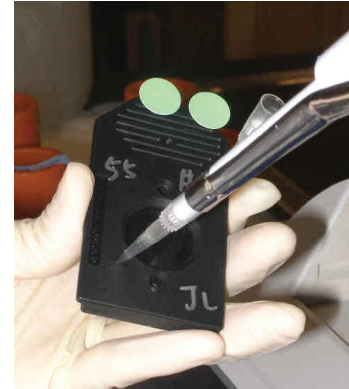
Hybridization Plate



7. Tightly seal the plate, vortex **TWICE** at high speed in 5 sector format, and spin down at 2000 rpm for 1 min.
8. Load the plate onto the thermal cycler and run the *CytoScan Hyb* program.

Temp	Time
95 °C	10 min
49 °C	Hold

9. Allow the samples to incubate at 49 °C for at least 1 min before loading.
10. Leaving the samples on the thermal cycler, load 200 µL of sample onto each array using a single-channel P200 pipette. Only hybridize up to 8 arrays at a time.



11. Clean any excess fluid from around the septa.
12. Apply Tough-Spots® to the septa and press firmly.
13. Immediately load the arrays into the hybridization oven, four at a time.
14. Hybridize the arrays 16 to 18 hrs at 50 °C and 60 rpm.

### IMPORTANT POINTS:

- Samples must remain on the thermal cycler while loading the arrays.
- To avoid damaging the septa, use a single-channel P200 pipette to load the arrays.
- If bubbles adhere to the array surface, tap the array lightly on the edge of a countertop, then gently shake the array a few times to ensure bubbles are not visible through the window.

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## CytoScan™ Assay 24 Samples: Stage 9 – Wash, Stain, Scan

1. Aliquot the following reagents into separate 1.5 ml microfuge tubes for each array:
  - A. 500  $\mu$ L Stain Buffer 1 into amber tubes
  - B. 500  $\mu$ L Stain Buffer 2 into clear tubes
  - C. 800  $\mu$ L Array Holding Buffer into blue tubes



500  $\mu$ L  
Stain Buffer 1



500  $\mu$ L  
Stain Buffer 2



800  $\mu$ L  
Array Holding Buffer

### Washing and Staining Arrays

1. Prime the Fluidics Station with the Wash buffers. Load the stain solutions and select the appropriate Fluidics Protocol.
  - If using the CytoScan™ HD Array, select "CytoScanHD\_Array\_450".
  - If using the CytoScan™ 750K Array, select "CytoScan750K\_Array\_450".
2. Start the Fluidics Protocol and leave the cartridge lever down in the Eject position.
3. Remove the Tough-Spots® from each array.
4. Load the arrays onto the Fluidics Station.

### Before Scanning

1. Ensure no bubbles are visible through the window.
2. Cover the septa with Tough-Spots®, then load onto the scanner.
3. Scan the arrays as described in the *CytoScan™ Assay User Guide* (PN 703038).

### Important Points

- Aliquot Stain1 Buffer into amber tubes.
- Aliquot Array Holding Buffer into blue tubes.
- Stain Buffer 1 and Array Holding Buffer are light sensitive.
- If there is a delay after aliquoting into the tubes, store the tubes at 4 °C, protected from light.
- Remove the bubbles from the arrays on the Fluidics Station (see the *Affymetrix® GeneChip® Fluidics Station 450/250 User's Guide*, P/N 08-0092) or remove the bubbles manually.

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