

# **FFPE-Derived DNA Quality Assessment**

## In Preparation for HaloPlex Target Enrichment

## Protocol

Version B0, July 2015

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#### **Manual Part Number**

G9900-90050

#### Edition

Version B0, July 2015

Printed in USA

Agilent Technologies, Inc. 5301 Stevens Creek Blvd Santa Clara, CA 95051 USA

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### In this Guide...

This guide describes a protocol for qualification of DNA extracted from formalin-fixed paraffin-embedded (FFPE) tissues for target enrichment using the Agilent HaloPlex system.

### 1 Before You Begin

This chapter contains information (such as required reagents and equipment) that you should read and understand before you start an experiment.

#### 2 Sample Preparation

This chapter describes the PCR-based DNA qualification protocol and includes guidelines for data analysis and suggested HaloPlex protocol modifications for FFPE-derived DNA samples.

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Make sure you read and understand the information in this chapter and have the necessary equipment and reagents listed before you start an experiment.



### **Required Reagents**

#### Table 1 Required Reagents for FFPE DNA Sample Anaysis

Description	Vendor and part number	
Reference DNA	need information	
Herculase II Fusion Enzyme with dNTPs (100 mM; 25 mM for each nucleotide), 200 reactions	Agilent p/n 600677	
Primer 105 FWD (hGAPDH Region 1–105 bp–Forward Primer): 5'-GGCTGAGAACGGGAAGCTTG-3'	General laboratory oligonucleotide supplier; HPLC purified	
Primer 105 REV (hGAPDH Region 1–105 bp–Reverse Primer): (5'-ATCCTAGTTGCCTCCCCAAA-3')	General laboratory oligonucleotide supplier; HPLC purified	
Primer 236 FWD (hGAPDH Region 2–236 bp–Forward Primer): (5'-CGGGTCTTTGCAGTCGTATG-3')	General laboratory oligonucleotide supplier; HPLC purified	
Primer 236 REV (hGAPDH Region 2 –236 bp– Reverse Primer:) (5'-GCGAAAGGAAAGAAAGCGTC-3')	General laboratory oligonucleotide supplier; HPLC purified	
10 mM Tris-HCl, pH 7.5–8.5, molecular biology grade	General laboratory supplier	
Nuclease-free Water (not DEPC-treated)	Ambion Cat #AM9930	
Quant-iT dsDNA BR Assay Kit, for use with the Qubit fluorometer		
100 assays, 2-1000 ng 500 assays, 2-1000 ng	Life Technologies p/n Q32850 Life Technologies p/n Q32853	
Agencourt AMPure XP Kit 5 mL 60 mL	Beckman Coulter Genomics p/n A63880 p/n A63881	

\* Also available separately as Herculase II Fusion DNA Polymerase, 40 reactions (Agilent p/n 600675) and 100 mM dNTP Mix (Agilent p/n 200415, sufficient for 1000 HaloPlex enrichment reactions).

### **Required Equipment**

Description	Vendor and part number	
2100 Bioanalyzer Platform and Consumables		
2100 Bioanalyzer Laptop Bundle	Agilent p/n G2943CA	
2100 Bioanalyzer Electrophoresis Set	Agilent p/n G2947CA	
High Sensitivity DNA Kit	Agilent p/n 5067-4626	
Thermal Cycler	Agilent SureCycler 8800, p/n G8800A or equivalent	
Thermal cycler-compatible 0.2-mL strip tubes or 96-well plates	See manufacturer's recommendations	
Magnetic separator compatible with strip tubes or 96-well plates	Agencourt SPRIPlate Super Magnet Plate p/n A32782, or equivalent	
Benchtop microcentrifuge	VWR p/n 93000-196, or equivalent	
P10, P20, and P200 pipettes	Pipetman P10, P20, P200, P1000 or equivalent	
Qubit 2.0 Fluorometer	Life Technologies p/n Q32866	
Qubit assay tubes	Life Technologies p/n Q32856	
Ice bucket	General laboratory supplier	
Vortex mixer	General laboratory supplier	

#### Table 2 Required Equipment for FFPE DNA Sample Anaysis

### **Safety Notes**

### CAUTION

• Wear appropriate personal protective equipment (PPE) when working in the laboratory.

### 1 Before You Begin

Safety Notes



## Protocol

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This section contains a PCR-based protocol for FFPE-derived DNA sample quality assessment and suggested modifications to the HaloPlex target enrichment protocol for FFPE-derived DNA samples.



### 2 Protocol Qualification of DNA Samples for HaloPlex Target Enrichment

### **Qualification of DNA Samples for HaloPlex Target Enrichment**

Results of target enrichment using the HaloPlex system are influenced by sample DNA integrity. Since DNA purified from formalin-fixed paraffin-embedded (FFPE) tissues can be highly degraded, it is useful to qualify the FFPE DNA samples prior to target enrichment in order to assess sample suitability for enrichment and to identify sequencing protocol modifications that may improve performance.

You can assess the suitability of your FFPE-derived DNA samples for HaloPlex target enrichment using the multiplex PCR-based qualification assay described here. In this protocol, each FFPE DNA sample is used as template for PCR amplification of two independent GAPDH amplicons. Yield of amplicons from the FFPE DNA template is measured and compared to yield of amplicons from an intact reference DNA template such as a HapMap DNA sample. The resulting sample-to-reference yield ratio serves as a quantitative indicator of DNA integrity which can be used to predict sample performance in HaloPlex target enrichment. Step 1. PCR-amplify GAPDH targets from FFPE-derived DNA and Reference DNA samples

# Step 1. PCR-amplify GAPDH targets from FFPE-derived DNA and Reference DNA samples

**1** Use the Qubit dsDNA BR Assay or PicoGreen staining kit to determine the concentration of your FFPE tissue-extracted DNA and the Reference DNA samples.

Follow the manufacturers instructions for the kits and instruments.

**2** Dilute the each DNA sample to a final DNA concentration of 5 ng/µL in 10 mM Tris-HCl, pH 7.5–8.5.

Store the DNA samples on ice.

- **3** Prepare each of the four hGAPDH PCR primers listed in Table 1 on page 8 at a final DNA concentration of  $24 \ \mu M$ .
- **4** Prepare the multiplex PCR master mix on ice by combining the reagents in the following table. Prepare enough master mix for the number of FFPE samples to be analyzed, plus one or more reference DNA samples, plus one reaction excess. Include the appropriate high-quality reference DNA sample(s) in each PCR amplification assay.

Reagent	Volume for 1 reaction	Volume for 12 reactions (includes excess)
Nuclease-free water	31.9 µL	414.7 μL
5X Herculase II Reaction Buffer	10 µL	130 µL
dNTPs (100 mM, 25 mM for each dNTP)	0.6 μL	7.8 μL
Primer 105 FWD (24 μM)	1.25 μL	16.25 μL
Primer 105 REV (24 μM)	1.25 μL	16.25 μL
Primer 236 FWD (24 µM)	1.25 μL	16.25 μL
Primer 236 REV (24 μM)	1.25 μL	16.25 μL
Herculase II Fusion DNA Polymerase	0.5 μL	6.5 μL
Total	48 µL	624 μL

**Table 3**Preparation of PCR master mix

5 Mix the master mix components by gentle vortexing, then distribute 48-µL aliquots to fresh 0.2-mL reaction tubes.

Step 1. PCR-amplify GAPDH targets from FFPE-derived DNA and Reference DNA samples

- **6** Add 2  $\mu$ L of the appropriate 5 ng/ $\mu$ L DNA sample to each tube.
- 7 Mix by gentle vortexing and then spin briefly to collect the liquid.
- 8 Place the amplification reaction tubes in a thermal cycler and run the program in Table 4, using a heated lid.

Segment	Number of Cycles	Temperature	Time
1	1	95°C	5 minutes
2	26	95°C	30 seconds
		60°C	60 seconds
		72°C	60 seconds
3	1	72°C	5 minutes
4	1	4°C	Hold

### Table 4 PCR program for FFPE DNA quality assessment

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### Step 2. Purify the amplicon DNA

In this step, the DNA amplicons are purified using AMPure XP beads.

- 1 Let the AMPure XP beads come to room temperature for at least 30 minutes.
- **2** Prepare 400 μL of 70% ethanol per sample, plus excess, for use in step 8.
- **3** Mix the AMPure XP bead suspension well, until the suspension appears homogeneous and consistent in color.
- 4 Add 100  $\mu$ L of the homogenous bead suspension prepared in step 3 to each 50- $\mu$ L PCR sample. Vortex thoroughly.

Using this bead-to-sample volume ratio is imperative to ensure optimal purification results.

**5** Incubate samples for 5 minutes at room temperature with continuous shaking.

Make sure the samples are properly mixing in the wells during the 5-minute incubation.

- **6** Spin briefly to collect the liquid, then place the tubes in the magnetic plate. Wait for the solution to clear (approximately 5 minutes).
- 7 Keep the tubes in the magnetic plate. Carefully remove and discard the cleared solution from each tube using a  $100-\mu L$  pipette set to  $100 \ \mu L$ . Do not touch the beads while removing the solution.
- 8 Continue to keep the tubes in the magnetic plate while you add 200  $\mu$ L of 70% ethanol into the tubes.

Use fresh 70% ethanol for optimal results.

- **9** Wait for 30 seconds to allow any disturbed beads to settle, then remove the ethanol using a 200-µL pipette set to 200 µL.
- **10** Repeat step 8 and step 9 once for a total of two washes.
- **11** Remove any residual ethanol with a  $20-\mu$ L volume pipette.
- **12** Air-dry the tubes with open lids at room temperature until the residual ethanol completely evaporates.

Make sure all ethanol has evaporated before continuing.

13 Remove tubes from the magnetic plate and add 40  $\mu$ L of 10 mM Tris-HCl to each sample.

#### 2 Protocol

Step 2. Purify the amplicon DNA

- 14 Mix thoroughly by pipetting up and down 15 times using a 100- $\mu$ L pipette set to 30  $\mu$ L.
- **15** Incubate for 2 minutes at room temperature to allow elution of DNA.
- **16** Put the tube in the magnetic plate and leave for 2 minutes or until the solution is clear.
- 17 Remove the cleared supernatant (approximately 40  $\mu L)$  to a fresh tube. You can discard the beads at this time.
- **Stopping Point** If you do not continue to the next step, samples may be stored at -20°C prior to analysis. Avoid subjecting the stored DNA samples to multiple freeze-thaw cycles.

# **Step 3. Measure amplicon yields from test and Reference DNA samples**

Use a Bioanalyzer High Sensitivity DNA Assay kit and the 2100 Bioanalyzer with 2100 Expert Software (version B.02.07 or higher required to run the High Sensitivity Kit). See the reagent kit guide for general Bioanalyzer instrument and assay setup instructions.

- 1 Prepare the chip, samples and ladder as instructed in the reagent kit guide, using 1  $\mu$ L of purified amplicons from the test sample and 1  $\mu$ L of purified amplicons from the Reference DNA sample for the analysis.
- **2** Load the prepared chip into the 2100 Bioanalyzer and start the run within five minutes after preparation.
- **3** Analyze the electropherogram for each sample.
  - Check that the electropherogram shows one peak with average size of 105 bp (±10%) and a second peak with average size of 236 bp (±10%).
  - Determine the concentration of both GAPDH amplicons (105-bp and 236-bp) in the sample by integration under each of the two peaks.

See Figure 1 for sample Bioanalyzer electropherograms.



**Figure 1** Example of 2100 Bioanalyzer system electropherograms for an intact reference DNA sample (left) and an FFPE-derived DNA sample (right).

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#### 2 Protocol

Step 4. Analyze amplicon yield ratio and identify protocol modifications

# Step 4. Analyze amplicon yield ratio and identify protocol modifications

Use the yield values determined in Step 3 to calculate the test-to-reference yield ratios for each amplicon. The avarage yield ratio may then be used as a quantitative measure of FFPE-derived DNA quality in order to predict performance in the HaloPlex target enrichment workflow. Suggested adjustments to the HaloPlex target enrichment protocol are provided.

**1** Calculate the yield ratio for the 105-bp amplicon according to the following formula:

105-bp ratio = 105-bp yield FFPE DNA/105-bp yield reference DNA

**2** Calculate the yield ratio for the 236-bp amplicon according to the following formula:

236-bp ratio = 236-bp yield FFPE DNA/236-bp yield reference DNA

- **3** Calculate the Average Yield Ratio by averaging the yield ratio values calulated for the 105-bp (from step 1) and 236-bp (from step 2) amplicons.
- **4** Use the guidelines in the table below to design modifications to the HaloPlex target enrichment protocol and subsequent sequencing protocol based on the apparent integrity of the FFPE-derived DNA sample. See "Typical Results" on page 19 for examples of post-enrichment Bioanalyzer profiles for FFPE-derived DNA samples of each category.

Sample Integrity Category	Average Yield Ratio	Recommended DNA Input (ng) in HaloPlex protocol	Recommeded additional sequencing
A	>0.2 (>20%)	200–500	1×–5×
В	0.05 to 0.2 (5% to 20%)	500–1000	5×–10×
С	<0.05 (<5%)	1000–2000	10×-100×

### NOTE

Intact DNA samples are enriched using the HaloPlex protocol at 200 ng of input DNA per reaction.

### **Typical Results**

Bioanalyzer profiles of the HaloPlex target-enriched libraries produced from different categories of FFPE-derived DNA samples are shown below. The lower abundance of long DNA fragments in FFPE-derived DNA samples results in deviations from the typical Bioanalyzer profile for intact DNA.



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### In This Book

This guide contains information to run the FFPE-Derived DNA Quality Assessment protocol in preparation for target enrichment using the HaloPlex system.

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Version B0, July 2015



G9900-90050

