

Title:

DNA analysis using Agilent High Sensitivity DNA Kit for 2100 Bioanalyzer Systems

Date last updated:

7th July 2024

Total Pages:

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Written by:

Jeanie Wu

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1. PURPOSE

To analyse DNA with sample sizes ranging from 50bp to 7000bp after NGS library preparation.

2. MATERIALS & EQUIPMENTS

Samples

- Dengue or Zika prepared NGS library

Reagents

- Agilent High Sensitivity DNA Kit, includes reagents and chips (Agilent, 5067-4626)

Consumables

- 0.2 ml PCR tubes
- Nuclease-free water

Equipments

- Agilent 2100 Bioanalyzer Systems (Bioanalyzer, chip priming station and IKA vortex mixer)
- Centrifuge (> 13000g)

3. PROCEDURE

Essential Measurement Practices

- Handle and store all reagents according to the instructions on the label of the individual box
- Avoid sources of dust or other contaminants. Foreign matter in reagents and samples or in the wells of the chip will interfere with assay results
- Keep all reagents and reagent mixes refrigerated at 2 – 8°C when not in use
- Allow all reagents and samples to equilibrate to room temperature for 30 min before use

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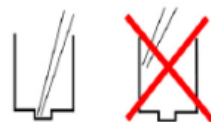
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- Protect dye and dye mixtures from light
- Always insert pipette tip to the bottom of the well when dispensing the liquid. Placing the pipette at the edge of the well may lead to poor results
- Use a new syringe and electrode cleaners with each new kit
- Use loaded chips within 5 min after preparation. Reagents might evaporate, leading to poor results
- Do not touch the 2100 Bioanalyzer instrument during analysis and never place it on a vibrating surface

**Step 1: Sample Preparation**

- For high sensitivity chip, prepare samples in the concentration range of 100pg/ul – 10ng/ul

Step 2: Setting up Chip Priming Station

- Replace the syringe by unscrewing old syringe from the lid of chip priming station
- Release old syringe from the clip and discard
- Remove plastic cap of the new syringe and insert into the clip
- Slide it into the hole of the luer lock adapter and screw it tightly to chip priming station
- Adjust the base plate by pulling the latch and opening the chip priming station
- Using a screwdriver, open the screw at the underside of the base plate
- Lift the base plate and insert it again in position C. Retighten the screw
- Adjust the syringe clip by releasing the lever of the clip and sliding it down to the lowest position



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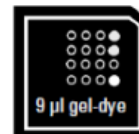
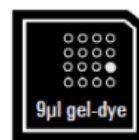
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Step 3: Preparing Gel-Dye Mix

- Allow High Sensitivity DNA dye concentrate (blue ●) and High Sensitivity DNA gel matrix (red ●) to equilibrate to room temperature for 30 min
- Add 15 ul of High Sensitivity DNA dye concentrate (blue ●) to a High Sensitivity DNA gel matrix vial (red ●)
- Vortex solution well and spin down. Transfer to spin filter
- Centrifuge at 2240 g \pm 20% for 15 min.
- Protect solution from light
- Store at 2 – 8°C. Use prepared gel-dye mix within 6 weeks of preparation

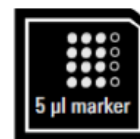
Step 4: Loading the Gel-Dye Mix

- Allow the gel-dye mix to equilibrate to room temperature for 30 min before use
- Put a new High Sensitivity DNA chip on the chip priming station
- Pipette 9 ul of gel-dye mix in the well marked 
- Make sure the plunger is positioned at 1 ml and then close the chip priming station
- Press plunger until it is held by the clip
- Wait for exactly 60 s then release clip
- Wait for 5 s, then slowly pull back the plunger to the 1 ml position
- Open the chip priming station and pipette 9 ul of gel-dye mix in the wells marked 



Step 5: Loading the Marker

- Pipette 5 ul of marker (green ●) in all sample and ladder wells. Do not leave any wells empty



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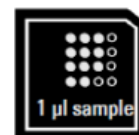
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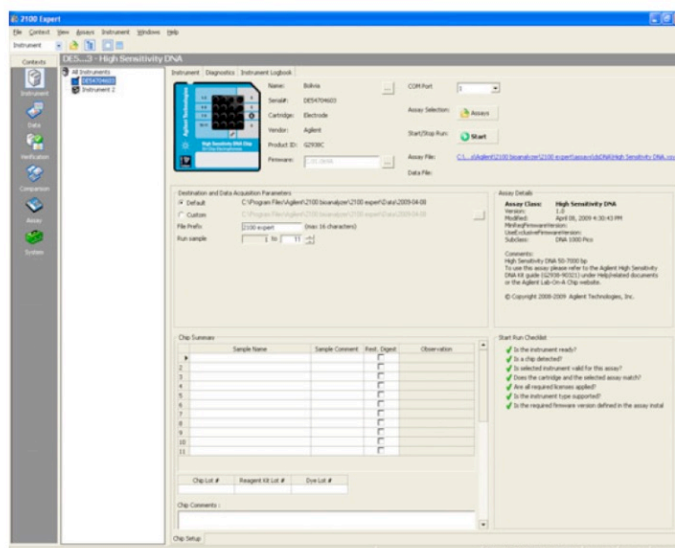
Step 6: Loading the Ladder and Samples

- Pipette 1 ul of High Sensitivity DNA ladder (yellow ) in the well marked 
- In each of the 11 sample wells, pipette 1 ul of sample (used wells) or 1 ul of marker (unused wells)
- Put the chip horizontally in the adaptor and vortex for 1 min at indicated setting (2400 rpm)
- Run the chip in the 2100 Bioanalyzer instrument within 5 min



Step 7: Inserting a Chip in the 2100 Bioanalyzer Instrument

- Open lid of 2100 Bioanalyzer instrument
- Place chip carefully into the receptacle. The chip fits only one way
- Carefully close the lid. The electrodes in the cartridge fit into the wells of the chip
- The 2100 Expert software screen shows that you have inserted a chip and upon closing the lid, a chip icon is displayed at the top left of the **Instrument** context



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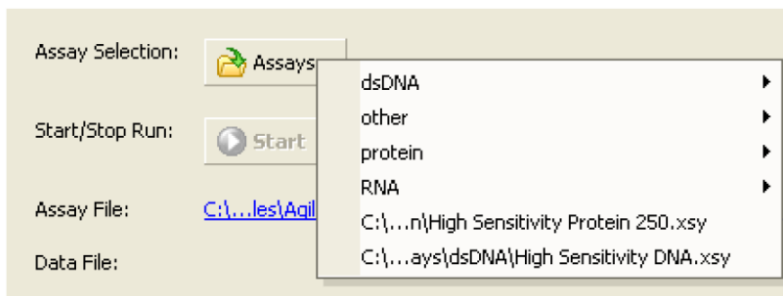
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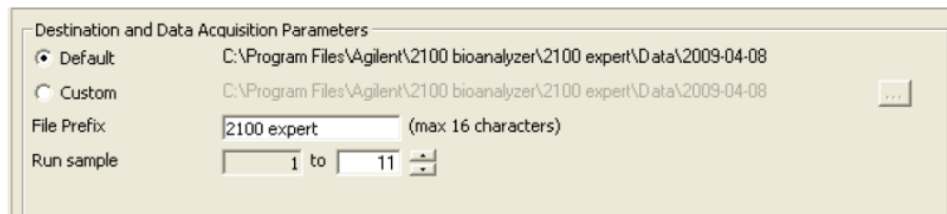
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Step 8: Starting the Chip Run

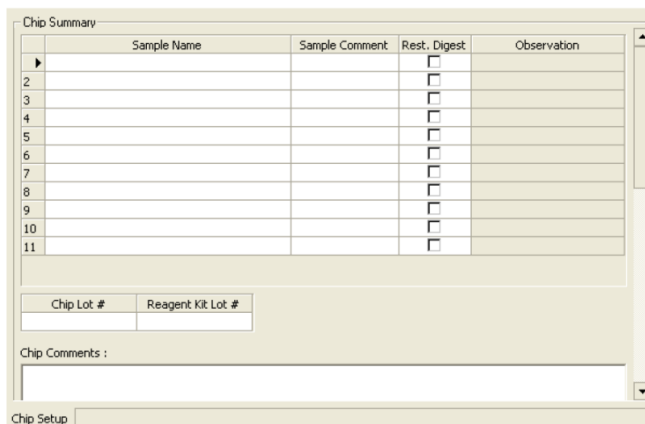
- a. In the **Instrument** context, select the appropriate assay from the “Assay” menu



- b. Data will be saved automatically to a file with a name using the prefix you have just entered. You can also customize the file storage location and the number of samples that will be analyzed.



- c. Complete sample name table by entering sample information like sample names and comments



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- d. Click the **Start** button in upper right of window to start the chip run. The incoming raw signals are displayed in the **Instrument** context.



- e. After the chip run is finished, remove the chip immediately and dispose it according to good laboratory practices.

Step 9: Cleaning Electrodes after a High Sensitivity DNA Chip Run

- a. Slowly fill one of the wells of electrode cleaner with 350 ul deionized analysi-grade water
- b. Open lid and place electrode cleaner in the 2100 Bioanalyzer instrument
- c. Close lid and leave it closed for about 10 – 30 s
- d. Open lid and remove electrode cleaner
- e. Wait another 30 s to allow water on the electrodes to evaporate before closing lid

High Sensitivity DNA ladder Well Results

To check results of your run, select the Gel or Electropherogram tab in the **Data** context. The electropherogram of the ladder well should resemble those shown below:

Major features of a successful ladder run are:

- 15 peaks for High Sensitivity DNA ladder
- All peaks are well resolved
- Flat baseline
- Correct identification of both markers

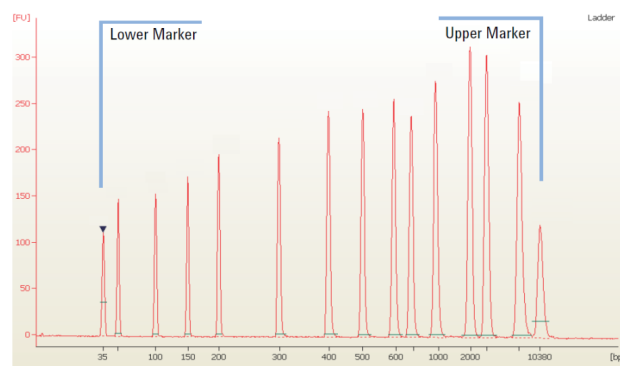


Figure 2 High Sensitivity DNA ladder

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High Sensitivity DNA Sample Well Results

To review the results of a specific sample, select the sample name in the tree view and highlight the *Results* sub-tab.

Typical Nextera libraries show a broad size distribution of ~ 250 – 1000 bp, as shown below.

Figure 1 Example Bioanalyzer Trace

