DS-02 Matrix Standard Kit (Dye Set E5)

SeqStudio™, 3500, 3730, and 3130 series instruments

Catalog Number 4323014

Pub. No. 4363121 Rev. E

WARNING! Read the Safety Data Sheets (SDSs) and follow the handling instructions. Wear appropriate protective eyewear, clothing, and gloves. Safety Data Sheets (SDSs) are available from **thermofisher.com/support**.

Product description

The DS-02 Matrix Standard Kit (Dye Set E5) is used to perform spectral calibrations when analyzing DNA fragments labeled with dR110™, dR6G™, dTAMRA™, dROX™, and LIZ™ dyes. (The LIZ™ dye is used to label the size standard.) The matrix standard contains five DNA fragments. Each fragment is labeled with a different dye from the dye set.

For more information on spectral calibration, see the *DNA Fragment Analysis by Capillary Electrophoresis User Guide* (Pub. No. 4474504).

Contents and storage

Contents	Amount	Storage
DS-02 Matrix Standard in TE buffer	1 tube	Store at 2–8°C, protected from light.[1] Do not freeze.

[1] The kit is stable for 1 year when stored at 2–8°C.

Guidelines for use

- For more information on the use of matrix standards, see the instrument user guide or getting started guide.
- To prepare the matrix standard dilution, combine the appropriate volumes of matrix standard and Hi-Di™ Formamide (Cat. No. 4311320). Dilution volumes vary depending on the instrument.
- Use the matrix standard within 2 hours of preparation.
- Do not add size standard to the matrix standard.
- Discard any unused reagent that has been diluted in Hi-Di™ Formamide.

Prepare the standard

1. Vortex the matrix standard tube for 5–10 seconds to mix, then centrifuge for 3–5 seconds to bring the mixture to the bottom of the tube and eliminate air bubbles.
2. Combine the volumes of matrix standard and Hi-Di™ Formamide (Cat. No. 4311320) appropriate for the instrument. See “Component volumes and well location for the prepared standard” on page 2.
3. Vortex for 5–10 seconds, then centrifuge for 3–5 seconds.
4. Dispense 10 µL of the prepared standard into the appropriate wells of a 96-well plate. See “Component volumes and well location for the prepared standard” on page 2.
5. Cover the plate with adhesive film, then centrifuge for 3–5 seconds.
6. Denature the DNA fragments:
   a. Incubate the mixture at 95°C for 5 minutes.
   b. Incubate the mixture at 4°C, or on ice, for ≥2 minutes.
7. Remove the adhesive film, then cover the plate with a 96-well septa (Cat. No. 4315933).
8. Centrifuge for 3–5 seconds.
9. Assemble the plate with the retainer and base, then load on the instrument.
10. Immediately perform the spectral calibration.

See the instrument user guide for specifics on setting up the run.

Component volumes and well location for the prepared standard

Table 1 SeqStudio™ Genetic Analyzer

Component	Volume	Well location for the prepared standard
	4-capillary array	
DS-02 Matrix Standard	1 µL	Dispense 10 µL of the prepared standard into wells of a 96-well plate: 4 wells (for example, A1–D1)
Hi-Di™ Formamide	49 µL	
Total volume	**50 µL**	

Table 2 3500/3500xL Genetic Analyzer

Component	Volume	Well location for the prepared standard
	8-capillary array 24-capillary array	
DS-02 Matrix Standard	3 µL	**Data Collection Software v3 and later:** Dispense 10 µL of the prepared standard into wells of a 96-well plate: • **8-capillary array**—8 wells (for example, A1–H1) • **24-capillary array**—24 wells (for example, A1–H3, A4–H6, A7–H9, or A10–H12) **Note:** If you place the standard in wells that do not correspond to injection position 1, specify the starting well position in the software. **Data Collection Software v1, v1.1, and v2:** Dispense 10 µL of the prepared standard into wells of a 96-well plate: • **8-capillary array**—8 wells: A1–H1 • **24-capillary array**—24 wells: A1–H3
Hi-Di™ Formamide	247 µL	
Total volume	**250 µL**	

Table 3 3730/3730xl DNA Analyzer

Component	Volume			Well location for the prepared standard
	48-capillary array		96-capillary array	
	Standard configuration	Reduced cross-talk (RCT) configuration[1]	Reduced cross-talk (RCT) configuration[1]	
DS-02 Matrix Standard	7 µL	13 µL	13 µL	Dispense 10 µL of the prepared standard into wells of a 96-well plate: • **48-capillary array**—48 wells (odd columns only): A1–H1, A3–H3, A5–H5, A7–H7, A9–H9, A11–H11 • **96-capillary array**—96 wells
Hi-Di™ Formamide	993 µL	987 µL	987 µL	
Total volume	**1,000 µL**	**1,000 µL**	**1,000 µL**	

[1] **For 3730/3730xl Data Collection Software only when running the RCT configuration:** Select dye set **Any5Dye-RCT** to perform fragment analysis in applications with a high dynamic range (large peaks with a signal intensity that is much higher than the signal intensity of small peaks).

Table 4 3130/3130*xl* Genetic Analyzer

Component	Volume	Well location for the prepared standard
	36-cm array 50-cm array	
DS-02 Matrix Standard	5 µL	Dispense 10 µL of the prepared standard into wells of a 96-well plate: • **16-capillary array**—16 wells: A1–H2 • **4-capillary array**—4 wells: A1–D1
Hi-Di™ Formamide	195 µL	
Total volume	**200 µL**	

Limited product warranty

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For descriptions of symbols on product labels or product documents, go to **thermofisher.com/symbols-definition**.

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Revision history: Pub. No. 4363121

Revision	Date	Description
E	10 April 2020	Correct error to the 3500 series instrument volumes introduced in Rev. C; revert to Rev. B volumes. Add vortex and centrifuge times. Add information for Data Collection Software v1, v1.1, and v2. Update format and licensing.
D	2 November 2018	Updated the compatible instruments and the manufacturing information.
C	18 August 2009	Baseline for this revision history

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