



Correct

## G. Run

- 1. Close the lid.
- 2. Press "Run Protocol"

full after loading.

40µl of sample into that well.

Make sure the the sample well

3. During a run, a marker peak should be detected for each cassette that is run. The marker runs ahead of the fractionation range.



# $\pmb{SageELF}^{{}^{\rm M}} \; {}^{\rm Quick} \; {}^{\rm Guide}$

0.75% Agarose Gel Cassette

for DNA fractionation between 10kb - 40kb

Product No.:	ELD4010
Cassette Description:	0.75% 10 kb - 40kb
Cassette Definition:	0.75% Agarose Marker 40
Software Version:	1.03

#### **Recommended Sample Load Guidelines**

**Ionic strength:** The ionic strength of the sample should be lower than the ionic strength of the buffer (80mM monovalent ions).

**Protein in the sample:** For best results, samples should be de-proteinized prior to loading.

Maximum Load: 5 µg Minimum Load: 100 ng

**Note:** this guide describes workflow using an internal marker. Refer to the SageELF operations manual for running timed fractionation.

## A. Prepare DNA samples

- 1. Bring DNA sample up to 30µl with TE.
- 2. Bring Loading Solution/Marker 40 mix to room temperature.
- 3. For each sample, combine 30µl of DNA sample with 10µl of loading solution/marker mix.
- 4. Mix samples thoroughly (vortex mixer). Briefly centrifuge to collect.

## **B.** Calibrate the Optics with the Calibration Fixtures

- 1. Place calibration fixture(s) onto the optical nest(s) as shown.
- 2. Close the instrument lid.
- 3. Go to the "Main" tab in the software.
- 4. Clear the protocol field by pressing the "Clear Run Data" button (if necessary).
- 5. Press the "Calibrate Detector" button.



- 6. In the Detector Calibration pop-up window, select the nest(s) to be calibrated.
- 7. Press "Start".
- 8. When calibration is complete, press "Return".

#### C. Program a Protocol

- 1. Go to the "Protocol Editor" tab in the software.
- 2. Select "New Protocol" in the menu bar.
- 3. Select 0.75% 10 kb-40kb from the "Casssette Definition" drop-down menu.
- 4. Using "size-based" mode, move the slider to a target elution well number.
- 5. Enter a value in to the "Target Value" window. This defines the range of fragments that will be collected in the target elution well. The collection range for the remaining wells will be calculated in the software.

			Separat	tion Mode e-based	Target Value,	bp Separati	on Time, hh:mm				<ul> <li>target value</li> </ul>
i	2	3	4	5	Target W	ell 7 8	9	10	ń		slider
	Estimated Size Distribution										
13678.1	10450.9	8388.0	6733.2	5232.8	4175.8	3300.1	2033.1	1615.7	1284.9	1022.3	🗲 target wells

6. The chart below can be used as a guideline to estimate fractionation values. .

Est. Run Time (hr)	1	2	3	4	5	6	7	8	9	10	11	12
6		28000	26500	19400	16862	13569	10872	8860	7646	6588	5000	4500
7		2660	28000	24000	17382	14595	12076	10085	8704	8562	5688	4500
7.5			43538	25474	21963	19076	16125	13544	10819	9452	6465	520
8			35175	28854	24388	20864	17857	15306	12645	10694	9300	650
8.5				39014	28658	23963	21338	18509	16278	13856	11530	980
9				62500	45506	25016	21864	19956	18390	16297	13667	1156
DNA compression occurs at this well												

7. Press "Save As" and name the fractionation protocol.

fingers

#### **D.** Prepare the Cassette(s)

1. While still taped closed! Hold the cassette sideways with the elution port side down. Tap to remove air bubbles from beneath the gel fingers.



- 2. Rotate the cassette in Rotate. the clock-wise direction. Allow the bubbles to collect in the lower buffer area. Gently tap if necessary. Allow bubbles to aggregate in lower buffer area. 3. Continue to rotate the Rotate. cassette clockwise, moving as many bubbles as possible to the side C E + S 9 L 8 6 0L LI ZI EL buffer area. Tap the Tap to dislodge cassette firmly to dislodge bubbles from any bubbles behind elution behind elution wells. wells. 4. Continue to rotate the cassette until most of the bubbles have accumulated in the upper buffer area. Place the cassette onto a benchtop. Continue to move the bubbles until most of the air is accumulated in the upper buffer area. 5. On a flat bench top: Peel off the adhesive tape. Grab the tape tab, hold the cassette firmly down, and pull the tape with a steady motion. 6. Remove all buffer from all 13 elution wells (set pipette to 40 µl to completely empty wells). Keep pipette tips vertical in the well to avoid damage to the membranes.
  - 7. Add 30 µl of buffer to all 13 elution wells.