MCE-202 MultiNA

APPLICATION 8

DNA-500

DNA-1000

DNA-2500

RNA

Accurate DNA Sizing Using the DNA-500 Reagent Kit

Using the MCE-202 MultiNA DNA-500 Reagent Kit, size analysis was conducted for a 25 bp to 100 bp sample and a 100 bp to 500 bp sample with accuracy of ±5 bp and ±5%, respectively.

K. Suzuki

Introduction

The MCE-202 MultiNA was used to generate a size calibration curve by ladder analysis, which in turn was used to conduct high-accuracy DNA size analysis. Here we introduce an example of size analysis using the DNA-500 Reagent Kit.

Results

Figure 1 shows the analysis results of the 25 bp DNA Step Ladder and pBR322-Hae III digest using the MCE-202 MultiNA DNA-500 Reagent Kit. For the 25 bp DNA Step Ladder (Fig. 1 (a)), 12 peaks from 25 bp to 300 bp were detected. In addition, a 1.8 kbp peak originating from the sample was detected in the vicinity of 116% on the migration time index. Analysis of the pBR322-Hae III digest (Fig. 1 (b)) yielded 17 peaks. In this experiment, the 123 bp and 124 bp fragments were recognized as a single peak, and no peaks were detected in the low weight density range of 8 bp to 21 bp. The size calculation results for the constituent fragments are shown in Table 1. The absolute error for the 25 bp to 100 bp fragments was less than 3 bp, and the relative error for the 100 bp to 500 bp fragments ranged from -0.9% to +3.4%.

(a) 25 bp DNA Step Ladder

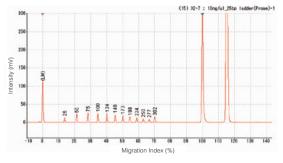


Table 1 DNA Sizing Results with the DNA-500 Reagent Kit

	Target Size Sizing Results (n = 8)					
		Average	RE		SD	RSD
	(bp)	(bp)	(%)	(bp)	(bp)	(%)
25bp step ladder	25	26.1	4.5	1.1	0.4	1.
	50	49.8	-0.5	-0.3	0.5	0.
	75	75.4	0.5	0.4	0.5	0.
	100	100.4	0.4	0.4	0.5	0.
	125	124.5	-0.4	-0.5	0.5	0.
	150	148.6	-0.9	-1.4	0.7	0.
	175	173.4	-0.9	-1.6	0.5	0
	200	198.8	-0.6	-1.3	0.9	0
	225	224.5	-0.2	-0.5	0.5	0.
	250	250.6	0.2	0.6	1.1	0
	275	277.0	0.7	2.0	0.9	0
	300	302.8	0.9	2.8	1.0	0
pBR322 – HaeIII digest	51	54.0	5.9	3.0	0.0	0
	57	57.9	1.5	0.9	0.4	0
	64	66.0	3.1	2.0	0.5	0
	80	82.4	3.0	2.4	0.5	0
	89	90.3	1.4	1.3	0.7	0
	104	107.5	3.4	3.5	0.9	0
	123	126.9	3.2	3.9	1.1	0
	184	186.6	1.4	2.6	1.5	0
	192	193.6	0.8	1.6	1.5	0
	213	215.6	1.2	2.6	1.4	0
	234	235.9	8.0	1.9	1.4	0
	267	267.9	0.3	0.9	1.0	0
	434	441.1	1.6	7.1	0.8	0
	458	467.1	2.0	9.1	1.1	0
	504	511.1	1.4	7.1	1.7	0
	540	548.1	1.5	8.1	1.2	0.
	587	603.5	2.8	16.5	2.4	0.

RE: Relative Error SD: Standard Deviation

RSD: Relative Standard Deviation (= SD / Average)

(b) pBR322-Hae III Digest

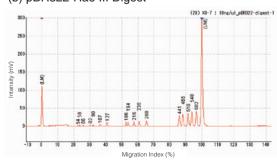


Fig. 1 Electropherogram of 25 bp DNA Step Ladder and pBR322-Hae III Digest Using the DNA-500 Reagent Kit for MultiNA

Analytical Conditions and Procedure

Instrument: MCE-202 MultiNA
Analysis mode: DNA-500 on-chip mixing

Ladder: 25 bp DNA ladder 20 ng/µL Sample: 25 bp DNA Step Ladder 10 ng/µL

pBR322-Hae III Digest 10 ng/µL

Reagents:

DNA-500 Reagent Kit for MultiNA

(Shimadzu) P/N 292-27910-91

 SYBR[®] Gold nucleic acid gel stain (Invitrogen)
 S-11494

25 bp DNA ladder

(Invitrogen) 10597-011

• 25 bp DNA Step Ladder

(Promega) G4511

pBR322 – Hae III Digest

(Sigma-Aldrich) D9655

Experimental Method:

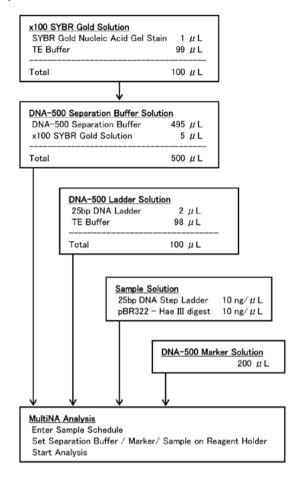


Fig. 2 Experimental Procedure (for 8 Samples)

(Note) For detailed operational information related to analysis using the MCE-202 MultiNA, please refer to the MCE-202 MultiNA Instruction Manual.



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