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ProductInformation

AGARASE I (EC 3.2.1.81)

Product Number A 8688 Lot Number 072K0729 Storage Temperature –20 °C

Product Description

SOURCE: Psuedomonas atlantica

Activity: 1 unit/ μl
DNase: None detected
RNase: None detected
Nickase: None detected
Protease: None detected

Functional Test: Suitable for use in isolating DNA

fragments from agarose gels.

Introduction

Agarase I is commonly used for isolating of DNA fragments of all sizes from agarose gels¹⁻⁴ prepared with low melting point agarose. This method³ is especially gentle and therefore useful for the isolation of very large DNA fragments (up to 680 kb) suitable for cloning pYAC libraries. This enzyme degrades melted agarose to oligosaccharides⁵ that will not gel on cooling. Agarase I and the oligosaccharides often do not interfere with subsequent manipulations, restriction endonuclease digestions, ligations or transformations for example. If necessary, the DNA can be separated from the agarase I reactants by routine phenol extraction and ethanol precipitation.²

Storage Buffer

40 mM Tris-HCl, pH 7.5 50 mM NaCl 50% glycerol

Assay Buffer

30 mM Bis-Tris, pH 7.1 at 25 °C 10 mM EDTA

Unit Definition

One unit is the amount of enzyme that digests 100 $\,\mu$ l of melted 1% low-melting agarose to neoagaro-oligosaccharides in 1 hour at 45 $\,^{\circ}$ C.

Usage

Cut the band of interest from the low melting point agarose (melting point 65 °C) gel. Place it in a tared reaction tube and determine the weight of the agarose slice (100 mg = 100 µl agarose gel). Add 0.04 volumes of 25X agarase I buffer (0.75 M Bis-Tris, 0.25 M EDTA, pH 7.1 at 25 °C). Incubate at 65 °C for 15 min. or until the agarose is melted. Cool the agarose to 45 °C. Add 1-2 units (1 unit for gels in 1X TAE, 2 units for gels in 1X TBE) agarase I per 100 mg of 1% agarose gel. Incubate for 1 hr at 45 °C. Note:Increase the agarase I proportionally for higher percentage agarose gels. Smaller DNA fragments (< 50 kb) can be removed from the reactants by ethanol precipitation as follows:

- Add 0.1 volume 3 M sodium acetate pH 5.2 (Product No. S 7899) to the melted agarose solution and incubate 15 minutes on ice.
- 2. Centrifuge 15 minutes to pellet the oligosaccharides.
- Remove the supernatant and place it in a suitable centrifuge tube and add 3 volumes of ice cold ethanol.
- 4. Centrifuge at 14,000 x g for 25 minutes and aspirate the ethanol.
- Air dry the DNA pellet and dissolve in DNase free water (Product No. W 4502), 1X TE (Product No. T 9285 diluted 1:100) or any other suitable buffer.

Functional Test

λDNA Hind III digest was run in a 1% agarose gel prepared from low melting agarose (Product No. A 9414) in 1X TBE (Product No. T 6400) running buffer. 6 bands (2.0-2.3kb) were excised and isolated as described in USAGE above. The isolated fragments were not degraded.

Activity Assay

Agarase is added to a reaction mixture containing assay buffer and melted agarose in 1 X TAE. The mixture is incubated 1 hour at 45 °C. The reaction is stopped by placing the reaction in an ice bath. After the tubes have cooled to 0 °C the reactions are tested for pipetablility. The least amount of enzyme that resulted in a pipetable reaction is defined as one unit.

DNase, RNase and Nickase

Nuclease assays are performed in assay buffer containing 5 units of agarase I and either λ -Hind III digest DNA, tRNA or pBR322 DNA and incubated 16 hours at 45 °C. The results were analyzed by agarose or polyacrylamide gel electrophoresis. No detectable DNase, RNase or nickase was found. Detection limit for DNase and RNase is > 10% degradation and for nickase is 1% conversion from supercoiled to linear or nicked circular DNA.

Protease

0.5% FITC-Casein was incubated with 10 μ l of enzyme in a 50 μ l reaction mixture for 1 hour at 37 °C. Liberated FITC equivalents are quantitated fluorometrically.

Detection limit: 8.5 X 10⁻⁶ μmoles of FITC released per minute.

References

- Burmeister M. F., and Lehrach H., Science 236, 1305 (1987).
- 2. Burmeister M., and Lehrach, H., Trends in Genetics, **5**, No 2, 41 (1989).
- 3. Imai, T. and Olson, M., Genomics, 8, 297 (1990).
- 4. Stefano, V., et al., Biochemistry International, **27**, 45 (1992).
- 5. Yaphe, W., Can. J. Microbiol., 3, 987 (1957).

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