

TGuide S32 Magnetic Tissue DNA Kit

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TGuide S32 Magnetic Tissue DNA Kit

Cat.no. 4992986

Kit Contents

Contents	4992986 (96 preps)
Buffer GHA	50 ml
Tissue DNA Reagents	6 plates
S32 Tip Comb	12 sets
Proteinase K	2 × 1 ml
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Animal tissue DNA reagent components

Column1/7	Column2/8	Column3/9	Column4/10	Column5/11	Column6/12
Buffer GHLp	Buffer GDAP	Buffer GDAP	Buffer PWDp	Buffer TB	MagAttract Suspension GSP1
600 µl	900 µl	900 µl	900 µl	100 µl	520 µl

Storage

This kit can be stored at room temperature (15-25°C) under dry condition for 12 months. For longer storage, we suggest storing the kit at 2-8°C. If a precipitate has formed in Buffer under 2-8°C, please place the buffer at room temperature or warm at 37°C for 10 min to dissolve the precipitate.

Introduction

The kit adopts magnetic beads with unique separation function and a unique buffer system to separate and purify high-quality genomic DNA from various animal tissues. The unique embedded magnetic beads have strong affinity for nucleic acid under certain conditions, and when the conditions change, the magnetic beads release adsorbed nucleic acid, thus achieving the purpose of fast separation and purification of nucleic acid.

The product is perfectly matched with TGuide S32 Automated Nucleic Acid Extractor. Magnetic beads are adsorbed, transferred and released by special magnetic rods, thus realizing the transfer of magnetic beads and nucleic acids. The whole experimental process is safe and convenient. The extracted genomic DNA fragments are large in size, with high purity, stable and reliable quality.

The DNA purified by the kit is suitable for various conventional operations, including enzyme digestion, PCR, library construction, Southern blot and other experiments.

Features

Easy and fast: Ultrapure genomic DNA can be obtained within 48 min.

Widely applicable: Applicable to all kinds of animal tissues.

Ultrapure: The obtained DNA has high purity and can be directly used in molecular biology experiments such as PCR, enzyme digestion, hybridization, etc.

Protocol

1. Preparation of DNA extraction reagents for animal tissues

Take out the vacuum package prepackaged 96-deep-well plate from the kit, mix it upside down for several times to resuspend the magnetic beads, remove the vacuum package. Gently shake the 96-deep-well plate to concentrate the reagent and magnetic beads to the bottom of the 96-deep-well plate (or centrifuge at 500 rpm for 1 min by the plate centrifuge). Carefully tear off the aluminum foil sealing film before use to avoid vibration of the 96-deep-well plate and prevent liquid spillage.

2. Sample treatment

Add 10-50 mg of tissue into 400 μ l of Buffer GHA and 20 μ l of Proteinase K, homogenize and centrifuge at 12,000 rpm for 1 min to remove impurities.

- 1) For samples with sufficient homogenate, the 65°C digestion step can be skipped.
- 2) For samples with visible tissue mass, it is recommended to digest at 65°C for 30 min to complete digestion.
- 3) For rat tail samples, perform the digestion at 56°C overnight.

Note: If RNA removal is required, add 4 μ l RnaseA (100 mg/ml) solution (self-provided), shake for 15 sec, and leave at room temperature for 5 min.

3. TGuide S32 Automated Nucleic Acid Extractor operation steps

- 3.1 Add 300 μ l of the above treated sample solution to columns 1 and 7 of the 96-well plate, and place the 96-deep-well plate on the 96-deep-well plate base of TGuide S32 Automated Nucleic Acid Extractor.
- 3.2 Insert the S32 Tip Combs into the slot of of magnetic rod tip comb slot of the TGuide S32 Automated Nucleic Acid Extractor.
- 3.3 Run the tissue automatic extraction program of TGuide S32 Automated Nucleic Acid Extractor.

Turn on the supporting Windows Pad, double-click the "Purification" icon to enter the TGuide S32 control program, click "Run", select the "DP602-Tissue" experimental program file and click the "Run" button in the lower right corner to start the experiment.

The specific experimental procedures are shown in the following table:

Step	Slot	Name	Waiting time (min)	Mixing time (min)	Adsorption time (sec)	Mixing speed	Volume (μ l)	Temperature ($^{\circ}$ C)	Strong adsorption mode
1	1	Lysis	0	2	0	Fast	900	--	--
2	6	Beads transfer	0	0.5	60	Fast	500	--	Yes
3	1	Beads adsorption	0	10	20	Fast	900	--	Yes
4	2	Washing 1	0	5	20	Fast	900	--	Yes
5	3	Washing 2	0	5	20	Fast	900	--	Yes
6	4	Washing 3	0	5	20	Fast	900	--	Yes
7	5	Elution	5	10	120	Fast	100	75	Yes
8	6	Beads disposal	0	0.5	0	Fast	500	--	--

- 3.4 Upon the completion of the automatic program, pipette out DNA sample in the 5th and 11th columns of the 96-deep-well plate and store under appropriate conditions.

Determination of DNA Concentration and Purity

The size of the obtained genomic DNA fragment is related to factors such as sample storage time and shearing force during operation. The concentration and purity of the obtained DNA fragments can be detected by agarose gel electrophoresis and ultraviolet spectrophotometer.

DNA should have a significant absorption peak at OD₂₆₀, with OD₂₆₀ value of 1 equivalent to about 50 μ g/ml double stranded DNA and 40 μ g/ml single stranded DNA.

The ratio of OD₂₆₀/OD₂₈₀ should be 1.7-1.9. If deionized water is used instead of elution buffer, the ratio will be low, because the pH value and the presence of ions will affect the light absorption value, but it does not mean the purity is low.