

ISOTISSUE

Code No. 317-03103 (For 3 Extractions), 311-03101 (For 30 Extractions)

Manual (ver.4) 20120106KH

I Product description

ISOTISSUE is a kit for extracting the genomic DNA from animal tissues. Since hazardous materials such as phenol and chloroform are not used during the handling and also a micro centrifuge tube is used, the handling is simple. The entire process can be completed in about 3 hrs. The genomic DNA obtained with this product is several 10 to several 100 kbp long and is highly purified so that it can be used as a substrate for restriction enzymes and for PCR.

II Precautions

- This product is a reagent for research and cannot be used for pharmaceutical or other purposes.
- We are not responsible for problems caused if this product is not handled in accordance with the manual.
- The product safety data sheet (SDS) can be reviewed on our website (www.nippongene.com).

III Contents of kit

Storage temperature: -20°C

	(3 extractions)	(30 extractions)
1. Extraction Buffer ^{(*)1}	1 ml × 2	20 ml
2. 20 mg/ml Proteinase K	150 µl	1.5 ml
3. 5 M NaCl	0.7 ml	7 ml
4. TE (pH 8.0)	1 ml × 2	20 ml
5. 10 mg/ml RNase A ^{(*)2}	20 µl	0.2 ml
6. 3 M Sodium Acetate (pH 5.2)	0.2 ml	1 ml × 2
7. 3 M LiCl ^{(*)3}	0.7 ml	7 ml
8. 150 mM Sodium Citrate ^{(*)3}	0.7 ml	7 ml

This product does not include isopropanol, ethanol or sterilized water.

(*)1 When SDS is deposited in the Extraction Buffer, melt the SDS by incubating at 37°C before use.

(*)2 White precipitates may be deposited in the RNase A solution but this does not indicate a quality or performance problem. In such cases, centrifuge and use only the supernatant.

(*)3 Prepare the Wash Solution just before use.

Wash Solution composition (final concentration):
0.3 M LiCl, 15 mM Sodium Citrate, 75% Ethanol

Preparation method (For 30 extractions):

Prepare by adding 3.5 ml of sterilized water and 52.5 ml of Ethanol (99.5%) to each of 7 ml solutions of 3 M LiCl and 150 mM Sodium Citrate included in the kit. The prepared solution will become cloudy. After preparing, store the solution at -20°C.

IV Examples of use

Sample (Tissue) 50 mg (1.5 ml microtube)

- ← Extraction Buffer 500 µl
Homogenize ^{(*)1}
- ← 20 mg/ml Proteinase K 40 µl
Vortex thoroughly
Incubate at 55°C for 15 min, while stirring once or twice ^{(*)2}
- ← 5 M NaCl 200 µl
Vortex for 1-2 sec.
(so that added NaCl becomes as homogeneous as possible)
Incubate at 55°C for 5 min ^{(*)3}
Leave standing on ice for 5 min (Precipitates containing SDS are formed)
Centrifuge (12,000×g, for 10 min, 4°C)

Supernatant ^{(*)4}

- ← Equal volume of isopropanol
Mix at room temperature for 5 min by gently inverting ^{(*)5}
Centrifuge (2,000×g, for 1 min)

Precipitates

- ← TE (pH 8.0) 400 µl
Dissolve by tapping ^{(*)6}
- ← Add 5 µl of 10 mg/ml RNase A, vortex for 1-2 sec.
Incubate at 55°C for 15 min.
- ← Add 5 µl of 20 mg/ml Proteinase K, vortex for 1-2 sec.
Incubate at 55°C for 15 min.
- ← Add 40 µl of 3 M Sodium Acetate (pH 5.2), vortex for 1-2 sec.
- ← Add 440 µl of isopropanol ^{(*)7}, vortex for 1-2 sec.
Centrifuge (2,000×g, for 1 min)

Precipitates

- ← Add 1 ml of the Wash Solution ^{(*)8} and wash ^{(*)9}
- ← Add 1 ml of the Wash Solution, and wash once more ^{(*)9} ^{(*)10}
- ← Wash ^{(*)9} with 1 ml of 70% ethanol.
- ← Wash ^{(*)9} with 1 ml of ethanol ^{(*)11}.
- ← Air dry ^{(*)12} for 5-10 min.
- ← Dissolve ^{(*)13} with 200-500 µl of TE (pH 8.0).

genomic DNA solution ^{(*)14}

*1) The tissue may be minced beforehand with sterilized scissors or the like or homogenized with a Polytron Homogenizer.

*2) If the tissue is not completely dissolved, extend the incubation time with Proteinase K by 15-20 min. However, avoid extending Proteinase K incubation time by 45 min or longer.

*3) When the solution cannot be homogeneous due to high viscosity, extend the incubation time for another 5 min.

*4) When the supernatant and the precipitates cannot be completely separated by centrifugation, withdraw only the supernatant so that the jelly-like lumps and tissue residues are not included. However, if separating the supernatant and the precipitates is impossible, it may be advisable to interrupt the extraction and switch to a method in which extraction is performed after homogenizing thoroughly in step *1).

*5) Mix thoroughly and as gently as possible so that the chromosomal DNA is not damaged.

*6) When the precipitates cannot be dissolved completely, incubate at 55°C for 15 min and gently tap the tube with a fingertip to dissolve. Further, if turbidity remains after dissolving, centrifuge (12,000 × g, 15 min), withdraw the supernatant so as not to collect the precipitates and use as a sample.

*7) In place of isopropanol, 2.5 volumes of ethanol can be used.

*8) See "III Kit content" (*3) for the preparation method of the Wash Solution.

*9) After mixing by inverting, carefully remove the supernatant using a Pipetman so as not to suck in DNA precipitates.

*10) If contamination by polysaccharides and the like is suspected, the frequency of washing by the Wash Solution may be increased further.

*11) To make drying easy, this step may be omitted.

*12) Do not dry using a vacuum drier. If the precipitates are completely dried, the solubility is drastically reduced.

*13) Almost completely solubilized by standing overnight at around 4°C. If not dissolved after standing at 4°C, incubate at 55°C. If not solubilized by this treatment, gently pipette using a pipette tip with a larger bore by cutting the tip end at 2-3 mm (high molecular weight (>100 kb) DNA is fragmented to 20 and several kb by vigorous pipetting).

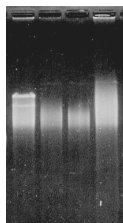
*14) Yield of genomic DNA obtained from mouse tissue

Mouse Tissue	DNA (μg/mg tissue)
Liver	0.5 ~ 1
Kidney	0.2 ~ 1
Heart	0.5 ~ 1.5

V Data

1) Genomic DNA obtained from mouse tissue

1 2 3 4

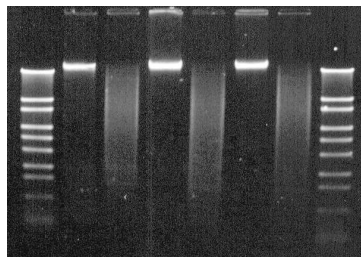


1. T4GT7 DNA : 166 kbp
2. Liver genomic DNA
3. Kidney genomic DNA
4. Heart genomic DNA

Pulse field gel electrophoresis

2) Restriction enzyme treatment of genomic DNA obtained from mouse tissue

1 2 3 4 5 6 7 8



1. Marker 6 (λDNA / Sty I)
2. Heart genomic DNA intact
3. Heart genomic DNA / EcoRI
4. Stomach genomic DNA intact
5. Stomach genomic DNA / EcoRI
6. Kidney genomic DNA intact
7. Kidney genomic DNA / EcoRI
8. Marker 6 (λDNA / Sty I)

Agarose S 0.8% gel, TAE Buffer 50V 80 min.

VI Troubleshooting

Q1. The sample is not dissolved or is difficult to dissolve.

A1. • Before adding the Extraction Buffer, mince the sample as finely as possible using sterilized scissors or the like.

• Homogenize the sample with a Polytron homogenizer or the like.

• Increase the volume of Proteinase K to be added.

• Use the sample after diluting by adding twice the normal volume of the Extraction Buffer. However, depending on the sample DNA may be degraded.

• Extend the incubation time of the Proteinase K treatment.

• The sample may be heat denatured. Lower the temperature of the Proteinase K treatment to an appropriate temperature and extend the reaction time (For example, 37°C for 3 hrs).

Q2. Yield is low.

A2. • If the sample is not completely dissolved, dissolve according to A1 above.

• If jelly-like material is found in the centrifuge fraction after the Proteinase K treatment, increase the volume ratio of the Extraction Buffer to the sample two-fold, homogenize and then extract.

Q3. OD₂₆₀ / OD₂₈₀ value is not more than 1.7.

A3. • Repeat washing DNA precipitates with the Wash Solution (0.3 M LiCl, 15 mM Sodium Citrate, 75% Ethanol).

• If no effect is observed, add to the DNA solution 300 mM LiCl at final concentration and 150 mM Sodium Citrate at final concentration. Incubate at 55°C for 15 min and then perform the ethanol precipitation and wash with the Wash Solution (0.3 M LiCl, 15 mM Sodium Citrate, 75% Ethanol).

Q4. DNA is not dissolved.

A4. • Reduce the frequency of homogenization.

• Increase the volume of the Extraction Buffer to be added.

• Perform gentle pipetting (Use a pipette tip with a larger bore by cutting the tip end at 2-3 mm).

Q5. DNA is broken down.

A5. • Avoid pipetting as much as possible.

• Perform pipetting as gently as possible (Use a pipette tip with a larger bore by cutting the tip end at 2-3 mm).

• Use fresh samples.

Q6. RNA is contaminated.

A6. • Redo, starting with the RNase treatment.