



2008

■ TISSUE SV MINI  
TISSUE SV MIDI  
TISSUE SV MAXI

exgene<sup>tm</sup>

**DNA PURIFICATION HANDBOOK**

  
GeneAll

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We thank you for any comment or advice.

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This protocol handbook is included in :

GeneAll® Exgene™ Tissue SV mini (104-101, 104-152)

GeneAll® Exgene™ Tissue SV Midi (104-226, 104-201)

GeneAll® Exgene™ Tissue SV MAXI (104-310, 104-326)

GeneAll® Exgene™ Tissue SV Plus! mini (109-101, 109-152)

GeneAll® Exgene™ Tissue SV Plus! Midi (109-226, 109-201)

GeneAll® Exgene™ Tissue SV Plus! MAXI (109-310, 109-326)

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## **Exgene™ Tissue SV**

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# Kit Contents

## GeneAll® Exgene™ Tissue SV (plus!) Series

| Cat. No.                  | 104(9)-101 | 104(9)-152 | 104(9)-226 |
|---------------------------|------------|------------|------------|
| Size                      | mini       | mini       | Midi       |
| No. of preparation        | 100        | 250        | 26         |
| Column type G             | 100        | 250        | 26         |
| Collection tube           | 300        | 750        | 52         |
| Buffer RL                 | 100 ml     | 250 ml     | 200 ml     |
| Buffer TL                 | 30 ml      | 80 ml      | 30 ml      |
| Buffer TB                 | 50 ml      | 110 ml     | 60 ml      |
| Buffer BW                 | 80 ml      | 220 ml     | 100 ml     |
| Buffer TW                 | 100 ml     | 250 ml     | 120 ml     |
| Buffer AE*                | 30 ml      | 60 ml      | 30 ml      |
| Proteinase K (20 mg/ml)** | 2.4 ml     | 6 ml       | 3.2 ml     |
| Protocol Handbook         | I          | I          | I          |

| Cat. No.                  | 104(9)-201 | 104(9)-310 | 104(9)-326 |
|---------------------------|------------|------------|------------|
| Size                      | Midi       | MAXI       | MAXI       |
| No. of preparation        | 100        | 10         | 26         |
| Column type G             | 100        | 10         | 26         |
| Collection tube           | 200        | 20         | 52         |
| Buffer RL                 | 250 ml x 3 | 200 ml     | 250 ml x 2 |
| Buffer TL                 | 120 ml     | 40 ml      | 100 ml     |
| Buffer TB                 | 240 ml     | 80 ml      | 240 ml     |
| Buffer BW                 | 400 ml     | 80 ml      | 220 ml     |
| Buffer TW                 | 250 ml x 2 | 120 ml     | 250 ml x 2 |
| Buffer AE*                | 120 ml     | 30 ml      | 60 ml      |
| Proteinase K (20 mg/ml)** | 12 ml      | 2.4 ml     | 6.4 ml     |
| Protocol Handbook         | I          | I          | I          |

\* 10mM Tris-HCl, pH 9.0, 0.5mM EDTA

\*\* After reconstitution of proteinase K, store it at 4 °C or -20 °C

## Storage Conditions

All components of GeneAll® Exgene™ Tissue SV kit should be stored at room temperature. After reconstitution of Proteinase K with storage buffer, it should be stored under 4 °C for conservation of activity. It can be stored at 4 °C for 1 year without significant decrease in activity. But for prolonged preservation of activity, storage under -20 °C is recommended.

During shipment or storage under cold ambient condition, a precipitate may formed in buffer TB. In such a case, heat the bottle at 37 °C to dissolve completely.

Otherwise, to use precipitated buffers will lead to poor DNA recovery. GeneAll® Exgene™ Tissue SV kits are guaranteed for 1 year.

## Quality Control

All components in GeneAll® Exgene™ Tissue SV kit are manufactured in strictly clean condition, and its degree of cleanness is monitored periodically. Restriction enzyme assay, PCR amplification assay and spectrophotometric assay as quality control are carried out from lot to lot thoroughly, and only the qualified is approved to be delivered.

## Safety Information

The buffers included in GeneAll® Exgene™ Tissue SV Kit contain irritant which is harmful when in contact with skin or eyes, or when inhaled or swallowed. Care should be taken during handling. Always wear gloves and eye protector, and follow standard safety precautions. Buffer TB contains chaotropes. It can form highly reactive compounds when combined with bleach.

Do NOT add bleach or acidic solutions directly to the sample-preparation waste.

# Product Specifications

## GeneAll® Exgene™ Tissue SV

|                        | mini       | Midi*      | MAXI*      |
|------------------------|------------|------------|------------|
| Sample size            | ~ 25 mg    | ~ 100 mg   | ~ 250 mg   |
| Preparation time       | 25 ~ min** | 40 ~ min** | 40 ~ min** |
| Maximum loading volume | 750 ul     | 5 ml       | 16 ml      |
| Binding capacity       | 60 ug      | 200 ug     | 600 ug     |
| Typical yield          | 20-60 ug   | 40-200 ug  | 100-600 ug |
| Elution volume         | 30-400 ul  | 200-600 ul | 0.4-2 ml   |

\* GeneAll® Exgene™ Tissue SV Midi/MAXI Kit procedures require the centrifuge which has a swining-bucket rotor and ability of 4,000 ~ 5,000 xg.

\*\* Depends on the complete lysis time of sample materials.

## Sample amount and expected yield

The amount of starting sample should not be exceed the recommended maximum limit, otherwise DNA recover will be significantly lowered. (Fig.1) Recommended amount of starting sample and the yield is listed on next page. If your starting material is not listed or you have no information about your sample, we recommend you start with smaller sample than the listed and increase the sample size in subsequent preparation depending on the result.

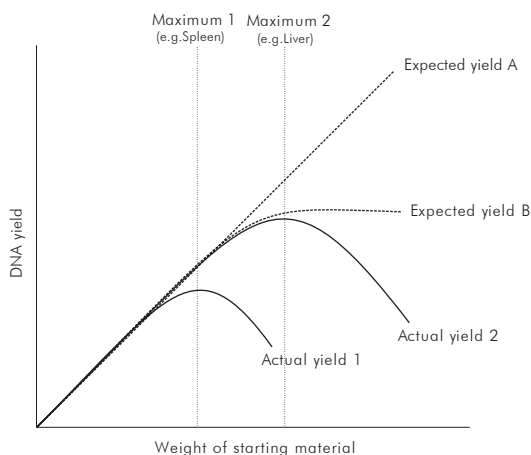


Fig. 1 Starting sample amount should not be exceed the maximum limit, otherwise DNA recover will be significantly lowered. If the cell mass of starting material is high (e.g., spleen, actual yield 1), maximum capacity will be lowered. (Maximum 1)

## Disruption and lysis

Generally, to make the sample finer will result in not only better yield and quality of DNA, but also reduced lysis time. Grinding in mortar and pestle under liquid nitrogen is a good method for disrupting the tissue sample, but alternative methods, such as a homogenizer or a bead-beater, can be employed in case by case. Shaking or vortexing during incubation for lysis may greatly accelerate the efficiency of lysis, followed by reduced time for complete lysis. Note that the freshness and the particle size of ground sample is the key for good result and that the fresh or frozen sample should be kept on ice until use.

**Table I. Maximum amount of starting sample**

| Sample           | Maximum amount      |                     |                      |
|------------------|---------------------|---------------------|----------------------|
|                  | mini                | Midi                | MAXI                 |
| Animal tissue    | 25 mg               | 100 mg              | 250 mg               |
| Mammalian blood* | 300 ul              | 2 ml                | 5 ml                 |
| Mouse tail       | 1 cm                | -                   | -                    |
| Rat tail         | 0.6 cm              | -                   | -                    |
| Insect           | 50 mg               | 200 mg              | 500 mg               |
| Cultured cell    | 5 x 10 <sup>6</sup> | 2 x 10 <sup>7</sup> | 5 x 10 <sup>7</sup>  |
| Bacteria         | 2 x 10 <sup>9</sup> | 8 x 10 <sup>9</sup> | 2 x 10 <sup>10</sup> |

\* Exgene™ Tissue SV *plus!* is required for whole blood

**Table 2. Typical DNA yields from various sample using Exgene™ Tissue SV mini.**

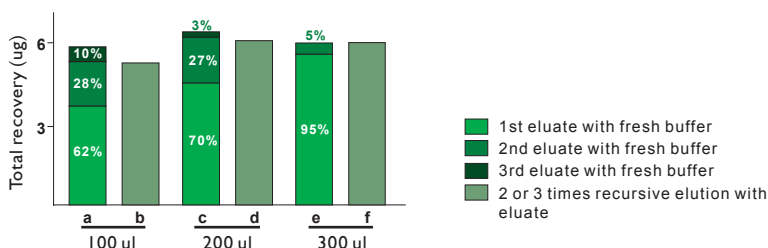
The yield of this table is calculated by addition of each eluate of 3 successive elution steps after DNA preparation with RNase A treatment. Without RNase A treatment, average yield from some sample may be significantly different from this data.

| Sample        | Starting amount     | Yield (ug) |
|---------------|---------------------|------------|
| Brain         | 20 mg               | 5-18       |
| Heart         | 20 mg               | 4-10       |
| Kidney        | 20 mg               | 15-35      |
| Liver         | 20 mg               | 15-35      |
| Lung          | 20 mg               | 4-10       |
| Pancreas      | 20 mg               | 8-25       |
| Spleen        | 10 mg               | 10-35      |
| Rat tail      | 0.6 cm              | 15-35      |
| Cultured cell | 2 x 10 <sup>6</sup> | 10-25      |
| Lymphocytes   | 5 x 10 <sup>6</sup> | 10-25      |
| Whole blood   | 300 ul              | 5-15       |
| Bacteria      | 2 x 10 <sup>9</sup> | 5-25       |

## Elution

Purified DNA can be eluted from SV column membrane in either deionized water or buffer AE which contains 0.5 mM EDTA and 10 mM Tris-HCl, pH9.0. Elution buffer should be equilibrated to room temperature before applying to SV column. Typically in mini kit, elution is carried out in two successive steps using 200  $\mu$ l Buffer AE each time. The volume of elution can be adjusted depending on the starting materials or the downstream applications, but it should be over the minimum requirements to wet the entire column membrane (50  $\mu$ l per column for mini) and should not be over 300  $\mu$ l. Basically, it is recommended for recovery of higher DNA concentration to decrease the elution volume to minimum, but total DNA recovery will decrease in this case. Otherwise, if maximum recovery is needed, the volume of elution buffer should be increased to elute as much as possible.

Unless the starting material has very low cell density, DNA bound to the SV column membrane may not be eluted completely with a single elution step. Approximately 60~85 % of DNA will be eluted in the first eluate, and the rest of bound DNA in the next. (Fig.2) However, a single elution with recommended volume of elution buffer will be sufficient to recover the amount of DNA required for multiple PCR reactions. For very small samples (containing less than 1  $\mu$ g of DNA), only a single elution in 50  $\mu$ l of Buffer AE or deionized water will be sufficient for complete elution of DNA.



**Fig. 2 The yield and concentration of purified DNA depending on the elution procedures.**

Total DNA was prepared from 10 mg of mouse liver using GeneAll® Exgene™ Tissue SV mini. Each preparation was exactly identical except the elution procedure; Elution was performed 3 times per column with 100  $\mu$ l (a) and 200  $\mu$ l (c), or 2 times per column with 300  $\mu$ l (e) of fresh Buffer AE. At the same time, another elution was carried out 3 times (b, d) or 2 times (f) by recursive use of eluate instead of fresh Buffer AE.

## Centrifuge in Midi/MAXI Kits

GeneAll® Exgene™ SV Midi and MAXI procedures require the conventional centrifuge which has a swinging-bucket rotor and ability of 4,000 ~ 5,000 xg. Use of fixed-angle rotor will cause inconsistent contact of SV column membrane with mixtures and/or buffers. Low g-force will lead to uncomplete removal of ethanol from SV column membrane and reduced volume of eluate. Usable centrifuges and rotors were listed below, but you can employ any equivalent.

| Company                                           | Centrifuge    | Rotor       |
|---------------------------------------------------|---------------|-------------|
| Beckman Coulter Inc.<br>(California, USA)         | Allegra X-15R | Sx4750      |
|                                                   | Allegra 25R   | Sx4750A     |
|                                                   |               | TS-5.1-500  |
| Eppendorf AG<br>(Hamburg, Germany)                | 5804/5804R    | A-4-44      |
|                                                   | 5810/5810R    |             |
| EYELA Inc.<br>(Tokyo, Japan)                      | 5800          | RS-410      |
|                                                   | 5900          | RS-410M     |
| Hanil Science Industrial Inc.<br>(Incheon, Korea) | Union 5KR     | R-WS1000-6B |
|                                                   | Union 55R     | W-WS750-6B  |
|                                                   | MF-550        | HSR-4S      |
|                                                   | HA1000-6      | WHSR-4S     |
|                                                   | HA1000-3      |             |
| Hettich AG<br>(Kirchlengern, Germany)             | Rotina 35     | 1717        |
|                                                   | Rotanta 460   | 1724        |
|                                                   | Rotixa 50S    | 5624        |

# GeneAll<sup>®</sup> Tissue and Tissue *plus!* SV mini

*\* Read this protocol carefully before experiment.*



## Protocol for Animal Tissue

### **Before experiment**

*Prepare the water bath to 56 °C.  
Equilibrate buffer AE to room temperature.  
All centrifugation should be performed at room temperature.  
Buffer TB may precipitate at cold ambient temperature.  
If so, completely dissolve it in 37 °C water bath.*

### **I. Disrupt up to 25 mg of tissue as described in step Ia, Ib or Ic, depending on the sample type.**

To disrupt the sample finer will accelerate lysis and reduce the lysis time.

For spleen tissue, up to 10 mg can be processed.

If the starting sample is larger than 25 mg (if spleen, 10 mg), increase the volume of buffer TL proportionally. For 50 mg of liver tissue, 400 µl of buffer TL is required.

For over 50 mg of tissue, Exgene<sup>™</sup> Tissue SV Midi or MAXI is recommended.

**Ia. For soft tissue, such as liver or brain, put up to 25 mg of the tissue into 1.5 ml tube, add 200 µl of Buffer TL, and homogenize thoroughly with microhomogenizer.**

**Ib. If microhomogenizer is not available or the tissue is not soft, grind the tissue to a fine powder with liquid nitrogen in a pre-chilled mortar and pestle. Put up to 25 mg of the powdered tissue into 1.5 ml tube. Add 200 µl of Buffer TL and pulse-vortex for 15 seconds.**

**Ic. If neither Ia nor Ib is available, mince up to 25 mg of tissue with sharp blade or scalpel as small as possible. Put the tissue into a 1.5 ml tube. Add 200 µl of Buffer TL and pulse-vortex for 15 seconds.**

**\*\*\* Alternatively, tissue samples can be effectively disrupted using some instruments, such as a rotor-stator homogenizer or a bead-beater.**

- 2. Add 20 ul of Proteinase K solution. Mix completely by vortexing or pipetting. Incubate at 56 °C until the sample is completely lysed.**

It is essential to mix the components completely for proper lysis.

If the sample amount is larger than 25 mg (if spleen, 10 mg), increase the amount of Proteinase K proportionally. For 50 mg of liver tissue, 40 ul of Proteinase K solution is required.

Lysis time varies from 10 min to 3 hr usually depending on the type of tissue processed and the disruption method (step1). The lysate should become translucent without any particles after complete lysis. Overnight lysis does not influence the preparation.

If the sample tube is incubated in water bath or heating block, vortex occasionally (2-3 times per hour) during incubation to lysis readily. *Lysis in shaking water bath, shaking incubator or agitator would be best for efficient lysis.*

- 3. Check!! If Buffer TB precipitates, pre-heat in a 37 °C water bath to dissolve completely.**

Precipitated buffer will cause significant decrease in recover yield.

- 4. Spin down the tube briefly to remove any drops from inside of the lid.**

- 5. Optional: If RNA-free DNA is required, add 4 ul of RNase solution (100 mg/ml, Cat.No. I17-960), vortex to mix thoroughly, and incubate for 2 min at room temperature.**

GeneAll® Exgene™ Tissue SV column has stronger affinity to DNA than RNA. Although the treatment of RNase is omitted, RNA occupies very small portion of eluates. RNA may inhibit some downstream enzymatic reactions, but will not inhibit PCR.

- 6. Add 400 ul of Buffer TB. Immediately vortex the tube to mix thoroughly. Spin down the tube briefly to remove any drops from inside of the lid.**

If the sample is larger than 25 mg (if spleen, 10 mg), increase the volume of buffer TB proportionally. For 50 mg of liver tissue, 800 ul of buffer TB is required.

- 7. Apply the mixture to the SV column. Centrifuge for 1 min at 6,000 xg above (>8,000 rpm). Replace the collection tube with new one (provided).**

If more than 25 mg (10 mg spleen) of tissue is processed, apply the mixture twice; apply 700  $\mu$ l of the mixture, spin down, discard the filtrate, re-insert to the empty collection tube, and repeat the step again until all of the mixture is applied to the SV column. If the mixture has not passed completely through the membrane, centrifuge again at full speed until all of the solution has been passed through.

Centrifuge at maximum speed will not affect the DNA recovery.

- 8. Add 600  $\mu$ l of Buffer BW. Centrifuge for 30 sec at 6,000 xg above (>8,000 rpm). Replace the collection tube with new one (provided).**

- 9. Apply 700  $\mu$ l of Buffer TW. Centrifuge for 30 sec at 6,000 xg above (>8,000 rpm). Discard the filtrate and reinsert the SV column back into the collection tube.**

- 10. Centrifuge at full speed (above 13,000 xg) for 1 min to remove residual wash buffer. Place the SV column in a fresh 1.5 ml tube (not provided).**

Care must be taken at this step for eliminating the carryover of buffer TW. If carryover of buffer TW occurs on column membrane, centrifuge again for 1 min at full speed before transferring to the new 1.5 ml tube

**I I. Add 200 ul of Buffer AE or sterilized water. Incubate for 2 min at room temperature. Centrifuge at full speed (> 13,000 xg) for 1 min.**

Ensure that the buffer AE or sterilized water is dispensed directly onto the center of SV column membrane for optimal elution of DNA.

Repeat of elution step with fresh 200 ul elution buffer will increase the total DNA yield significantly, while a third elution step with a further 200 ul of elution buffer will increase yields slightly. Each eluate can be separated in fresh tubes or can be collected to same tube, but more than 300 ul of eluate can not be collected in a 1.5 ml tube because the SV column will come into contact with the eluate.

If higher concentration of DNA is needed or starting sample amount is very small, second elution can be carried out with the first eluate instead of the fresh elution buffer. Alternatively or simultaneously, elution volume can be decreased to 50 ul for higher DNA concentration. However the small volume of elution buffer will decrease the total yield of DNA recovery.

For long-term storage, eluting in Buffer AE is recommended. But EDTA included in Buffer AE may inhibit subsequent enzymatic reactions, so you can avoid such latent problem by using distilled deionized water (>pH7.0) or Tris-Cl (>pH8.5).

When using water for elution, check the pH (>7.0) of water before elution.



## Protocol for Rodent Tail

### **Before experiment**

Prepare the water bath to 56

Equilibrate buffer AE to room temperature

All centrifugation should be performed at room temperature.

*Buffer TB may precipitate at cold ambient temperature.*

*If so, dissolve it in 37 °C water bath.*

Prepare Ice and 0.5M EDTA solution (pH 8.0)

- 1. Add 30 ul of a 0.5M EDTA solution (pH 8.0) to 180 ul of Buffer TL in a 1.5 ml centrifuge tube. Chill on ice before use.**

- 2. Mince 0.3 - 1.0 cm of mouse or rat tail as small as possible. Transfer it to the tube containing chilled EDTA-Buffer TL mixture.**

For rodent tail tip, up to 0.6 cm (rat or adult mouse) or 1.0 cm (mouse) can be processed for each prep.

The tail should be submerged in EDTA-Buffer TL mixture.

**\*\*\* Alternatively, tissue samples can be effectively disrupted using some instruments, such as a rotor-stator homogenizer or a bead-beater.**

- 3. Add 20 ul of Proteinase K solution (20 mg/ml, provided).**

- 4 Incubate at 56 °C until the tissue is completely lysed.**

It can take up to 8 hours to lysis completely.

Vortex occasionally during incubation, or incubate on shaking or agitating platform.

Make sure the tail is completely digested.

Overnight lysis does not affect DNA recovery.

Complete lysate may appear clear and viscous. If the lysate appears to be gelatinous or has leftover particles, increase the lysis time or the volume of EDTA-Buffer TL and proteinase K. Remember that the volume of Buffer TB should be increased proportionally in subsequent step.

- 5. Continue with step 3 of Animal Tissue protocol (page 12).**



## Protocol for Cultured Animal Cell or Lymphocytes

### **Before experiment**

*Prepare the water bath to 56 °C.*

*Equilibrate buffer AE to room temperature.*

*All centrifugation should be performed at room temperature.*

*Buffer TB may precipitate at cold ambient temperature.*

*If so, completely dissolve it in 37 °C water bath.*

\* Washing cells with sufficient volume of PBS before procedures usually brings about better results.

### **1. Harvest cells (up to $5 \times 10^6$ cells) to a 1.5 ml microcentrifuge tube by centrifugation at 14,000 xg for 20 sec.**

Alternatively, cells can be pelleted at 1,000 xg for 5 min. For adherent cells, trypsinize the cells before harvesting.

Certain cells, such as PC12, do not lyse well in Buffer TL. For those cells, it is helpful to perform additional freeze-thaw step several times before proceeding to next step.

### **2. Discard the supernatant as much as possible and thoroughly resuspend cell pellet in 200 $\mu$ l of Buffer TL.**

For efficient resuspending in buffer TL, it is helpful to loosen and resuspend the pellets by flickering or vortexing before the addition of buffer TL

### **3. Add 20 $\mu$ l of Proteinase K solution. Mix completely by vortexing, or pipetting. Incubate for 10 min at 56 °C.**

Longer incubation will not affect DNA recovery.

### **4. Continue with Animal Tissue Protocols from step 3 (page 12).**



## Protocol for Insects or Worms

### **Before experiment**

Prepare the water bath to 56 °C.

Equilibrate buffer AE to room temperature.

All centrifugation should be performed at room temperature.

Buffer TB may precipitate at cold ambient temperature.

If so, completely dissolve it in 37 °C water bath.

- 1. Grind up to 50 mg (25 mg, if worms) of insect in liquid nitrogen with pre-chilled mortar and pestle. Place the powder in a 1.5 ml microcentrifuge tube.**

Fine powder will reduce lysis time and bring about better result.

Worms can be minced using sharp blade or scalpel instead of grinding in liquid nitrogen.

\*\*\* *Alternatively, tissue samples can be effectively disrupted using some instruments, such as a rotor-stator homogenizer or a bead-beater.*

- 2. Apply 200 ul of Buffer TL and mix completely by vortexing or pipetting.**
- 3. Continue with Animal Tissue Protocols from step 2 (page 12).**



## Protocol for Paraffin-fixed Tissue

### *Before experiment*

*Prepare xylene, absolute ethanol*

*Prepare the water bath to 56 °C.*

*Equilibrate buffer AE to room temperature*

*All centrifugation should be performed at room temperature.*

*Buffer TB may precipitate at cold ambient temperature.*

*If so, completely dissolve it in 37 °C water bath.*

### **1. Place a small section of paraffin-fixed tissue in a 2 ml microcentrifuge tube.**

Start with smaller sample. If DNA yield is smaller than expected, increase the amount of sample gradually in next preparations.

### **2. Add 1200 µl xylene. Vortex vigorously.**

### **3. Centrifuge at full speed (> 13,000 xg) for 5 min. Carefully remove supernatant by pipetting.**

Do not remove any of the pellet.

### **4. Add 1200 µl of absolute ethanol to the pellet to remove residual xylene and mix gently by vortexing.**

### **5. Centrifuge at full speed for 5 min. Carefully remove the ethanol by pipetting.**

Do not remove any of the pellet.

### **6. Repeat steps 4-5 for 2 times or more.**

### **7. Evaporate the ethanol by incubating the microcentrifuge tube at room temperature for 10-15 min with open cap.**

### **8. Resuspend the tissue pellet in 200 µl Buffer TL and follow the tissue protocol from step 2 (page 12).**



## Protocol for Gram Negative Bacteria

### Before experiment

Prepare the water bath to 56 °C.

Equilibrate buffer AE to room temperature.

All centrifugation should be performed at room temperature.

Buffer TB may precipitate at cold ambient temperature.

If so, completely dissolve it in 37 °C water bath.

### 1. Pellet bacterial cells (up to $2 \times 10^9$ cells) to a 1.5 ml microcentrifuge tube by centrifugation at 14,000 xg for 30 sec.

1~2 ml of overnight bacterial liquid culture ( $OD_{600} = 1$ ) may correspond to  $1 \times 10^9$  cells.

### 2. Discard the supernatant as much as possible and thoroughly resuspend bacterial pellet in 200 ul of Buffer TL.

For efficient resuspending in buffer TL, it is helpful to loosen and resuspend the pellets by flickering or vortexing before the addition of buffer TL

### 3. Add 20 ul of Proteinase K solution. Mix completely by vortexing or pipetting. Incubate for 10 min at 56 °C.

After complete lysis, lysis mixture will turn to clear from turbid. If the 10-min lysate still looks turbid or cloudy, incubate until the lysate become clear without any particle.

Lysis time may vary depending on the species and cell numbers. Cells can be further incubated for complete lysis and longer incubation time does not affect recover yield. After incubation, cool the lysate to room temperature.

### 4. Continue with Animal Tissue Protocols from step 3 (page 12).



## Protocol for Mammalian Whole Blood (*plus! only*)

\* For convenient preparation from whole blood, use GeneAll® Exgene™ Blood SV kit.

### **Before experiment**

Prepare the water bath to 56 °C.

Equilibrate buffer AE to room temperature.

All centrifugation should be performed at room temperature.

*Buffer TB may precipitate at cold ambient temperature.*

*If so, completely dissolve it in 37 °C water bath.*

### **1. Transfer 300 ul of whole blood to a new 1.5 ml tube.**

Before transfer of blood, it is recommended that gently rocking the tube of blood until thoroughly mixed.

### **2. Add 900 ul of Buffer RL to the tube containing the blood sample. Invert the tube 5-6 times to mix. Incubate the mixture for 10 min at room temperature.**

Invert 2-3 times during the incubation. The lysate should become clear (translucent). If the lysate is turbid (opaque) not clear, it may be frozen or mis-stored sample, and you should read the annotation of step 3.

If fresh or well-stored sample is processed, it will take less times than 10 min to acquire translucent mixtures.

For larger blood sample, we recommend GeneAll® Exgene™ Blood SV Midi/MAXI kit. For simple and convenient preparation from blood samples, we recommend GeneAll® Exgene™ Blood SV mini (105-101, 105-152). This kit provides the rapid and easy method for the purification of total DNA from up to 400 ul of whole blood or its derivative in addition to lymphocytes and cultured cells.

### **3. Centrifuge for 1 min at 14,000 xg (full speed) and carefully remove the supernatant as much as possible without disturbing the visible white (or pink) pellet. Resuspend the pellet by vortexing or flickering.**

Approximately 10-20 ul of residual liquid will remain. Steps 2-3 are critical steps for DNA recovery, so you have to check the translucent lysate and the white(or pink) pellet before processing next step. If blood sample has been frozen or mis-stored, resuspend the pellet and repeat step 2-3 with the resuspended cells until the lysate becomes translucent.

- 4. Check!! If Buffer TB precipitates, pre-heat in a 37 °C water bath to dissolve completely.**

Precipitated buffer will cause significant decrease in recover yield.

- 5. Add 200 µl of Buffer TL to the tube containing the resuspended cells. Pipet the solution 5-6 times to mix well.**

The lysate may be viscous.

- 6. Add 10 µl of Proteinase K solution. Mix thoroughly by vortexing or pipetting. Incubate for 10 min at 56 °C.**

Overnight lysis is available and it will not influence the preparation.

- 7. Continue with Animal Tissue Protocols from step 4 (page 12).**

# GeneAll<sup>®</sup> Exgene<sup>™</sup> Tissue SV Midi

\* Visit [www.geneall.com](http://www.geneall.com) for more Midi protocols.



## Protocol for Animal Tissue

### Before experiment

Prepare the water bath to 64 .  
Equilibrate buffer AE to room temperature.  
All centrifugation should be performed at room temperature.  
Buffer TB may precipitate at cold ambient temperature.  
If so, completely dissolve it in 37 water bath.

### Important Note!

All centrifugation steps **MUST** be carried out on a tabletop centrifuge which has an ability of 4,000 xg at least and a swing-out bucket.

DO NOT USE A FIXED-ANGLE ROTOR.

### I. Disrupt 30 ~ 100 mg of tissue as described in step Ia, Ib or Ic, depending on the sample type.

Finer sample powder will accelerate lysis and decrease the lysis time.

For spleen tissue, up to 40 mg can be processed. For over 100 mg of tissue, GeneAll<sup>®</sup> Exgene<sup>™</sup> Tissue SV MAXI (up to 250 mg) is recommended.

#### Ia. For soft tissue, such as liver or brain, put up to 100 mg of the tissue into 15 ml conical tube, add 400 ul of Buffer TL, homogenize thoroughly with microhomogenizer, add 600 ul of Buffer TL, and vortex vigorously to resuspend well.

#### Ib. If microhomogenizer is not available or the tissue is not soft, grind the tissue to a fine powder with liquid nitrogen in a pre-chilled mortar and pestle. Put up to 100 mg of the powdered tissue into 15 ml conical tube. Add 1 ml of Buffer TL and pulse-vortex for 30 sec.

#### Ic. If neither Ia nor Ib is available, mince the tissue with sharp blade or scalpel as small as possible. Put up to 100 mg of the tissue into a 15 ml conical tube. Add 1 ml of Buffer TL and pulse-vortex for 30 sec.

\*\*\* Alternatively, tissue samples can be effectively disrupted using some instruments, such as a rotor-stator homogenizer or a bead-beater.

- 2. Add 100 ul of Proteinase K solution to the tube. Mix completely by vortexing or pipetting. Incubate at 64 °C until the sample is completely lysed.**

It is essential to mix the components completely for proper lysis.

Lysis time varies from 10 min to 3 hr usually depending on the type of tissue processed and the disruption method (step1). The lysate should become translucent without any particles after complete lysis. Overnight lysis does not influence the preparation.

If the sample tube is incubated in water bath or heating block, vortex occasionally (2-3 times per hour) during incubation to lysis readily. Incubation in shaking water bath, shaking incubator, or agitator would be best for efficient sample lysis.

- 3. Check!! If Buffer TB precipitates, pre-heat in a 37 °C water bath to dissolve completely.**

Precipitated buffer will cause significant decrease in recover yield.

- 4. Optional: If RNA-free DNA is required, add 20 ul of RNase solution (100 mg/ml, Cat.No. I17-960), vortex to mix thoroughly, and incubate for 3 min at room temperature.**

GeneAll® Exgene™ Tissue SV column has stronger affinity to DNA than RNA. Although the treatment of RNase is omitted, RNA occupies very small portion of eluates. RNA may inhibit some downstream enzymatic reactions, but will not inhibit PCR.

- 5. Add 2 ml of Buffer TB. Immediately vortex or invert the tube to mix thoroughly.**

Check that buffer TB is not precipitated.

For efficient binding of DNA to membrane, it is essential to mix the sample thoroughly to yield a homogeneous solution.

- 6. Transfer the sample mixture to the SV Midi column carefully, close the cap, centrifuge for 2 min at 3,000 xg (3,800 rpm).**

There may be appear some floating matters in mixture.

It does not affect DNA recovery, so it is recommended transfer of even that matters.

If the mixture has not passed completely through the membrane, centrifuge again at higher speed until all of the solution has passed through.

- 7. Discard the filtrate and re-insert the SV column back into the 15 ml tube. Apply 3 ml of Buffer BW and centrifuge for 2 min at 3,000 xg (3,800 rpm).**

While transfer of mixture to the SV column, be careful not to moisten the rim of SV column.

- 8. Discard the filtrate, wipe off any spillage from the thread of the 15 ml tube, and re-insert the SV column back into the 15 ml tube.**

- 9. Carefully, without moistening of the rim, apply 4 ml of buffer TW, and centrifuge for 15 min at 4,500 xg (5,000 rpm).**

At least, 4,000 xg is required for proper DNA recovery.

Insufficient centrifugal force may lead to remaining of ethanol in SV column membrane, followed by poor DNA recovery.

If the SV column has buffer TW associated with it after centrifugation, incubate the SV column for 10~15 min at room temperature to evaporate residual ethanol. Residual ethanol may and inhibit some downstream reactions, such as PCR.

- 10. Place the SV column into a new 15 ml centrifugation tube (provided). Pipet 300 ul of buffer AE or distilled water onto a center of membrane and close the cap. Incubate for 2 min at room temperature. Centrifuge at full speed (over 4,000 xg, at least) for 5 min.**

Lower centrifugal force will dramatically reduce the volume of eluate.

Before this elution step, it is strongly recommended that any residual ethanol originated from buffer TW should not remain in SV column membrane. It can be checked by smell. Residual ethanol disturbs proper DNA preparation as follows; poor DNA recovery, low purity, inhibition of subsequent reactions and etc.

Ensure that the Buffer AE or distilled water is dispensed directly onto the center of the SV column membrane for optimal elution of DNA.

For blood sample, if starting volume is less than 1 ml, apply 200 ul of buffer AE or distilled water. Do not reduce the elution volume below 100 ul.

For long-term storage, eluting in buffer AE is recommended. But, EDTA included in the buffer AE may inhibit subsequent enzymatic reactions, so you can avoid such latent problems by using distilled water (>pH7.0) or Tris-Cl (>pH8.5). When using water for elution, make sure the pH of water (at least 7.0) before elution.

- 11. For higher concentrated yield, re-load the eluate from step 11 into the SV column, close the cap, incubate 2 min at room temperature, and centrifuge at full speed for 5 min.**

**For higher total yield, add 300 ul of fresh Buffer AE or distilled water again into the SV column, close the cap, incubate 2 min at room temperature, and centrifuge at full speed for 5 min.**

Less than 300 ul of eluate will be obtained from 300 ul of elution buffer, but this has no influence on DNA yields.



## Protocol for Cells

\* Visit [www.geneall.com](http://www.geneall.com) for more Midi protocols.

### Before experiment

Prepare the water bath to 64 °C.

Equilibrate buffer AE to room temperature.

All centrifugation should be performed at room temperature.

Buffer TB may precipitate at cold ambient temperature.

If so, completely dissolve it in 37 °C water bath.

### Important Note!

All centrifugation steps **MUST** be carried out on a tabletop centrifuge which has an ability of 4,000 xg at least and a swing-out bucket.

DO NOT USE A FIXED-ANGLE ROTOR.

- 1. Harvest cells listed below to a 15 ml conical tube by centrifugation at 2,000 xg for 10 min.**

Cultured cells or lymphocytes: up to  $2 \times 10^7$

Bacterial cells: up to  $8 \times 10^9$

- 2. Discard the supernatant as much as possible and thoroughly resuspend cell pellet in 1 ml of Buffer TL.**

For efficient resuspending in buffer TL, it is helpful to loosen and resuspend the pellets by flickering or vortexing before the addition of buffer TL.

- 3. Add 100 µl of Proteinase K solution. Mix completely by vortexing or pipetting. Incubate for 15 min at 64 °C.**

Longer incubation will not affect DNA recovery.

- 4. Continue with Animal Tissue Protocols from step 3 (page 23).**



# Protocol for Mammalian Whole Blood (*plus!* only)

\* For convenient preparation from whole blood, use GeneAll® Exgene™ Blood SV kit.

## Before experiment

Prepare the water bath to 64 °C.

Equilibrate buffer AE to room temperature

All centrifugation should be performed at room temperature.

Buffer TB may precipitate at cold ambient temperature.

If so, dissolve it in 37 °C water bath.

## Important Note!

All centrifugation steps **MUST** be carried out on a tabletop centrifuge which has an ability of 4,000 xg at least and a swing-out bucket.

DO NOT USE A FIXED-ANGLE ROTOR.

### 1. Transfer 2 ml of whole blood to a new 15 ml conical tube.

Before transfer of blood, it is recommended that gently rocking the tube of blood until thoroughly mixed.

### 2. Add 6 ml of Buffer RL to the tube containing the blood sample. Invert the tube 5-6 times to mix. Incubate the mixture for 10 min at room temperature.

Invert 2-3 times during the incubation. The lysate should become clear (translucent). If the lysate is turbid (opaque) not clear, it may be frozen or mis-stored sample, and you should read the annotation of step 3.

For simple and convenient preparation from blood samples, we recommend GeneAll® Exgene™ Blood SV.

For larger blood sample, we recommend GeneAll® Exgene™ Tissue SV *plus!* MAXI

### 3. Centrifuge for 10 min at 2,000 xg and carefully remove the supernatant as much as possible without disturbing the visible white (or pink) pellet. Resuspend the pellet by vortexing or flickering.

A little of residual liquid will remain. Steps 2-3 are critical steps for DNA recovery yields, so you have to check the translucent lysate and the white (or pink) pellet before processing next steps.

If blood sample has been frozen or mis-stored, resuspend the pellet and repeat step 2-3 with the sample until the lysate becomes translucent.

- 4. Add 1 ml of Buffer TL to the tube containing the resuspended cells. Pipet the solution 5-6 times to mix well.**

The lysate may be viscous.

- 5. Add 60 ul of Proteinase K solution. Mix thoroughly by vortexing. Incubate for 15 min at 64 .**

- 6. Check!! If Buffer TB precipitates, pre-heat in a 37 water bath to dissolve completely.**

Precipitated buffer will cause significant decrease in recover yield.

- 7. Continue with Animal Tissue Protocols from step 4 (page 23).**



## Protocol for Animal Tissue

### Before experiment

Prepare the water bath to 64 .  
Equilibrate buffer AE to room temperature.  
All centrifugation should be performed at room temperature.  
*Buffer TB may precipitate at cold ambient temperature.  
If so, completely dissolve it in 37 water bath.*

### Important Note!

*All centrifugation steps MUST be carried out on a tabletop centrifuge which has an ability of 4,000 xg at least and a swing-out bucket.*

DO NOT USE A FIXED-ANGLE ROTOR.

### I. Disrupt 100 ~ 250 mg of tissue as described in step Ia, Ib or Ic, depending on the sample type.

To make the sample finer will accelerate lysis and decrease the lysis time.

For spleen tissue, up to 100 mg can be processed.

#### Ia. Disrupt tissue sample using a hand-held homogenizer or a rotor-stator homogenizer in 3 ml of Buffer TL. Be careful not to overflow due to foaming.

#### Ib. If any homogenizer is not available, grind tissue sample to a fine powder with liquid nitrogen in a pre-chilled mortar and pestle. Put up to 250 mg of the powdered tissue into 15 ml conical tube. Add 3 ml of Buffer TL and pulse-vortex for 30 sec.

#### Ic. If neither Ia nor Ib is available, mince the tissue with sharp blade or scalpel as small as possible. Put up to 250 mg of the tissue into a 15 ml conical tube. Add 3 ml of Buffer TL and pulse-vortex for 30 sec.

\*\*\* *Alternatively, tissue samples can be effectively disrupted using some instruments, such as a bead-beater.*

- 2. Add 200 ul of Proteinase K solution to the tube. Mix completely by vortexing or pipetting. Incubate at 64 until the sample is completely lysed.**

It is essential to mix the components completely for proper lysis.

Lysis time varies from 10 min to 3 hr depending on the type of tissue processed and the disruption method (step1). The lysate should become translucent without any particles after complete lysis. Overnight lysis does not influence the preparation.

If the sample tube is incubated in water bath or heating block, vortex occasionally (3-4 times per hour) during incubation to lysis readily. Incubation in shaking water bath, shaking incubator, or agitator would be best for efficient sample lysis.

- 3. Check!! If Buffer TB precipitates, pre-heat in a 37 water bath to dissolve completely.**

Precipitated buffer will cause significant decrease in recover yield.

- 4. Optional: If RNA-free DNA is required, add 200 ul of RNase solution (100 mg/ml, Cat.No.117-961), vortex to mix thoroughly, and incubate for 3 min at room temperature.**

GeneAll® Exgene™ Tissue SV column has the very stronger affinity to DNA than RNA. Although the treatment of RNase is omitted, RNA occupies very small portion of eluates. RNA may inhibit some downstream enzymatic reactions, but will not inhibit PCR.

- 5. Add 7 ml of Buffer TB. Immediately vortex or invert the tube to mix thoroughly.**

Check that buffer TB is not precipitated.

For efficient binding of DNA to membrane, it is essential to mix the sample thoroughly to yield a homogeneous solution.

- 6. Transfer the sample mixture to the SV MAXI column carefully, close the cap, centrifuge for 2 min at 2,000 xg (3,000 rpm).**

There may be appear some floating matters in mixture. It does not affect DNA recovery, so it is recommended to transfer even that matters.

If the mixture has not passed completely through the membrane, centrifuge again at higher speed until all of the solution has passed through.

- 7. Discard the filtrate and re-insert the SV column back into the 50 ml tube. Apply 7 ml of Buffer BW and centrifuge for 2 min at 2,000 xg (3,000 rpm).**

While transfer of mixture to the SV column, be careful not to moisten the rim of SV column.

- 8. Discard the filtrate, wipe off any spillage from the thread of the 50 ml tube, and re-insert the SV column back into the 50 ml tube.**

- 9. Carefully, without moistening of the rim, apply 10 ml of buffer TW, and centrifuge for 15 min at 4,500 xg (5,000 rpm).**

At least, 4,000 xg is required for proper DNA recovery.

Insufficient centrifugal force may lead to remaining of ethanol in SV column membrane, followed by poor DNA recovery.

If the SV column has buffer TW associated with it after centrifugation, incubate the SV column for 10~15 min at room temperature to evaporate residual ethanol from buffer TW. Residual ethanol can inhibit some downstream reactions, such as PCR.

**10. Place the SV column into a new 50 ml conical tube (provided).**

**Pipet 600 ul of buffer AE or distilled water onto a center of membrane and close the cap. Incubate for 2 min at room temperature.**

**Centrifuge at 4,500 xg (5,000 rpm) for 5 min.**

Lower centrifugal force will dramatically reduce the volume of eluate.

Before this elution step, it is strongly recommended that any residual ethanol originated from buffer TW should not remain in SV column membrane. It can be checked by smell. Residual ethanol disturbs proper DNA preparation as follows; poor DNA recovery, low purity, inhibition of subsequent reactions and etc.

Ensure that the Buffer AE or distilled water is dispensed directly onto the center of the SV column membrane for optimal elution of DNA.

Do not reduce the elution volume below 400 ul.

For long-term storage, eluting in buffer AE is recommended. But, EDTA included in the buffer AE may inhibit subsequent enzymatic reactions, so you can avoid such latent problems by using distilled water (>pH7.0) or Tris-Cl (>pH8.5). When using water for elution, make sure the pH of water (at least 7.0) before elution.

**11. For higher concentrated yield, re-load the eluate from step 10 into the SV column, close the cap, incubate 2 min at room temperature, and centrifuge at 4,500 xg (5,000 rpm) for 5 min.**

**For higher total yield, add 600 ul of fresh Buffer AE or distilled water again into the SV column, close the cap, incubate 2 min at room temperature, and centrifuge at 4,500 xg (5,000 rpm) for 5 min.**

Less than 600 ul of eluate will be obtained from 600 ul of elution buffer, but this has no influence on DNA yields.



## Protocol for Cells

\* Visit [www.geneall.com](http://www.geneall.com) for more MAXI protocols.

### Before experiment

Prepare the water bath to 64 °C.

Equilibrate buffer AE to room temperature.

All centrifugation should be performed at room temperature.

Buffer TB may precipitate at cold ambient temperature.

If so, completely dissolve it in 37 °C water bath.

### Important Note!

All centrifugation steps **MUST** be carried out on a tabletop centrifuge which has an ability of 4,000 xg at least and a swing-out bucket.

DO NOT USE A FIXED-ANGLE ROTOR.

### 1. Harvest cells listed below to a 15 ml conical tube by centrifugation at 2,000 xg for 15 min.

Cultured cells or lymphocytes: up to  $5 \times 10^7$

Bacterial cells: up to  $2 \times 10^{10}$

### 2. Discard the supernatant as much as possible and thoroughly resuspend cell pellet in 3 ml of Buffer TL.

For efficient resuspending in buffer TL, it is helpful to loosen and resuspend the pellets by flickering or vortexing before the addition of buffer TL

### 3. Add 200 ul of Proteinase K solution. Mix completely by vortexing or pipetting. Incubate for 20 min at 64 °C.

Longer incubation will not affect DNA recovery.

### 4. Continue with Animal Tissue Protocols from step 3 (page 30).



## Protocol for Mammalian Whole blood(plus! only)

\* For convenient preparation from whole blood, use GeneAll® Exgene™ Blood SV kit.

### Before experiment

Prepare the water bath to 64 °C.

Equilibrate buffer AE to room temperature

All centrifugation should be performed at room temperature.

Buffer TB may precipitate at cold ambient temperature.

If so, dissolve it in 37 °C water bath.

### Important Note!

All centrifugation steps **MUST** be carried out on a tabletop centrifuge which has an ability of 4,000 xg at least and a swing-out bucket.

DO NOT USE A FIXED-ANGLE ROTOR.

### 1. Transfer 5 ml of whole blood to a new 50 ml conical tube.

Before transfer of blood, it is recommended that gently rocking the tube of blood until thoroughly mixed.

### 2. Add 15 ml of Buffer RL to the tube containing the blood sample. Invert the tube 5-6 times to mix. Incubate the mixture for 10 min at room temperature.

Invert 2-3 times during the incubation. The lysate should become clear (translucent). If the lysate is turbid (opaque) not clear, it may be frozen or mis-stored sample, and you should read the annotation of step 3.

For simple and convenient preparation from blood samples, we recommend GeneAll® Exgene™ Blood SV.

### 3. Centrifuge for 15 min at 2,000 xg and carefully remove the supernatant as much as possible without disturbing the visible white (or pink) pellet. Resuspend the pellet by vortexing or flickering.

A little of residual liquid will remain. Steps 2-3 are critical steps for DNA recovery yields, so you have to check the translucent lysate and the white (or pink) pellet before processing next steps.

If blood sample has been frozen or mis-stored, resuspend the pellet and repeat step 2-3 with the sample until the lysate becomes translucent.

- 4. Add 3 ml of Buffer TL to the tube containing the resuspended cells. Pipet the solution 5-6 times to mix well.**

The lysate may be viscous.

- 5. Add 150 ul of Proteinase K solution. Mix thoroughly by vortexing. Incubate for 20 min at 64 .**

- 6. Continue with Animal Tissue Protocols from step 3 (page 30).**



# Troubleshooting Guide for Exgene™ Tissue SV

| Facts              | Possible Causes                                                                                            | Suggestions                                                                                                                                                                                                                                                                                                                 |
|--------------------|------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Low or no recovery | Too much starting material                                                                                 | Too much starting material brings about inefficient lysis and/or SV column clogging, followed by poor DNA yields. Reduce the amount of starting material.                                                                                                                                                                   |
|                    | Starting material is too old or mis-stored                                                                 | Best results can be obtained with fresh sample. DNA yield is dependent on the type, size, age and storage of starting material. Lower yields will be obtained from material that has been inappropriately stored.<br>For example, blood samples that have been stored at 4 °C for more than 5 days may give reduced yields. |
|                    | Low cells in the sample                                                                                    | Low cell-density of starting sample leads to poor result. Increase the sample amount and load the SV column several times. Reduce the elution volume to 50 µl.<br>If possible, harvest fresh sample and repeat the DNA purification with a new sample.                                                                      |
|                    | Cell pellet was not resuspended thoroughly in step 2 of cultured cell protocol or step 3 of Blood protocol | For proper cell lysis with Buffer TL, it is essential to resuspend thoroughly the cell pellet. Inefficient cell lysis leads to many problems including poor DNA yields.                                                                                                                                                     |
|                    | Improper centrifuge (Midi/MAXI)                                                                            | Swing-bucket rotor should be used instead of fixed angle rotor.                                                                                                                                                                                                                                                             |
|                    | G-force in the protocol was not reach to 4,000 xg (Midi/MAXI)                                              | For proper DNA purification, centrifugal g-force in washing step ought to reach 4,000 xg at least (page 10).                                                                                                                                                                                                                |

| <b>Facts</b>                              | <b>Possible Causes</b>                                               | <b>Suggestions</b>                                                                                                                                                                                                                                       |
|-------------------------------------------|----------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b>Low or no recovery</b>                 | Insufficient lysis                                                   | Incomplete lysis is due to too much starting material, imperfect mixing with Buffer TL, insufficient time to lysis completely or poor disruption of sample. In next purification, check it carefully.                                                    |
|                                           | Weaken activity of proteinase K caused by mis-storage or out-of-date | Proteinase K must be stored under 4 °C for maintenance of proper activity. However, it is recommended to store at -20 °C for prolonged preservation of its activity. Lysis can not be done properly with degenerated proteinase K. Replace with new one. |
|                                           | Improper eluent                                                      | As user's requirement, elution buffer other than Buffer AE can be used. However, the conditions of optimal elution should be low salt concentration with alkaline pH (7<pH<9). When water or other buffer was used as eluent, ensure that condition.     |
| <b>Low A<sub>260/280</sub> ratio</b>      | Insufficient lysis                                                   | Insufficient lysis causes low DNA purity, and is usually due to too much starting material used, imperfect mixing with Buffer TL, insufficient time to lysis completely, or poor disruption of sample. Check these out in next preparations.             |
| <b>Low concentration of DNA in eluate</b> | Low cells in starting material or small starting material used       | Increase the amount of starting material with additional volume of buffer. Or, reduce the elution volume to the minimum and/or do re-elution with eluate.                                                                                                |
| <b>Column clogging</b>                    | Insufficient lysis                                                   | Insufficient lysis may lead to column clogging. In next preparations, mix the sample with each buffer completely, reduce the starting materials, extend the incubation time at 56 °C (or 65 °C), or increase the amount of proteinase K to double.       |

| <b>Facts</b>                                                      | <b>Possible Causes</b>                            | <b>Suggestions</b>                                                                                                                                                                                                        |
|-------------------------------------------------------------------|---------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b>White precipitate in Buffer TB</b>                             | Buffers were stored in cool ambient condition     | Storage at low temperature may cause precipitation in buffer TB. For proper DNA purification, any precipitate in the buffer should be dissolved completely by incubating the buffer at 37 (or above) until it disappears. |
| <b>Enzymatic reaction is not performed well with purified DNA</b> | High salt concentration in eluate                 | Ensure that all washing steps were carried out just in accordance with the protocols. Additional washing helps usually remove high salt in eluate. Refer the annotation of elution step.                                  |
|                                                                   | Low purity of DNA                                 | Check " Low $A_{260}/_{280}$ ratio "                                                                                                                                                                                      |
| <b>Degraded DNA</b>                                               | Starting material is too old or mis-stored        | Too old or mis-stored sample often yield degraded DNA. Use fresh sample.                                                                                                                                                  |
| <b>DNA floats out of well while loading of agarose gel</b>        | Residual ethanol from buffer TW remains in eluate | Ensure that wash step in protocols is performed properly. SV column membrane should be completely dried via additional centrifugation or air-drying.                                                                      |

# Ordering Information

※ For more information about ordering, visit [www.geneall.com](http://www.geneall.com)

| Products                                                    | Cat. No.                      | Type               | Size               |
|-------------------------------------------------------------|-------------------------------|--------------------|--------------------|
| <b>GeneAll® Exprep™</b> for the preparation of plasmid DNA  |                               |                    |                    |
| Plasmid SV mini                                             | 101-150<br>101-102<br>101-111 | mini/ spin/ vacuum | 50<br>200<br>1,000 |
| Plasmid SV Midi**                                           | 101-226<br>101-250<br>101-201 | Midi/ spin/ vacuum | 26<br>50<br>100    |
| Plasmid Quick                                               | 101-050<br>101-002<br>101-011 | mini/ spin         | 50<br>200<br>1,000 |
| <b>GeneAll® Expin™</b> for the purification of fragment DNA |                               |                    |                    |
| Gel SV                                                      | 102-150<br>102-102            | mini/ spin/ vacuum | 50<br>200          |
| PCR SV                                                      | 103-150<br>103-102            | mini/ spin/ vacuum | 50<br>200          |
| CleanUp SV                                                  | 113-150<br>113-102            | mini/ spin/ vacuum | 50<br>200          |
| Combo GP                                                    | 112-150<br>112-102            | mini/ spin/ vacuum | 50<br>200          |
| <b>GeneAll® Exgene™</b> for the extraction of total DNA     |                               |                    |                    |
| Tissue SV mini (plus!)*                                     | 104(9)-101<br>104(9)-152      | mini/ spin/ vacuum | 100<br>250         |
| Tissue SV Midi (plus!)**                                    | 104(9)-226<br>104(9)-201      | Midi/ spin/ vacuum | 26<br>100          |
| Tissue SV MAXI (plus!)**                                    | 104(9)-310<br>104(9)-326      | MAXI/ spin/ vacuum | 10<br>26           |
| Blood SV mini                                               | 105-101<br>105-152            | mini/ spin/ vacuum | 100<br>250         |
| Blood SV Midi**                                             | 105-226<br>105-201            | Midi/ spin/ vacuum | 26<br>100          |
| Blood SV MAXI**                                             | 105-310<br>105-326            | MAXI/ spin/ vacuum | 10<br>26           |

| Products                                                       | Cat. No. | Type               | Size   |
|----------------------------------------------------------------|----------|--------------------|--------|
| <b>GeneAll® Exgene™</b> <i>for the extraction of total DNA</i> |          |                    |        |
| Cell SV mini                                                   | 106-101  | mini/ spin/ vacuum | 100    |
|                                                                | 106-152  |                    | 250    |
| Cell SV MAXI**                                                 | 106-310  | MAXI/ spin/ vacuum | 10     |
|                                                                | 106-326  |                    | 26     |
| Clinic SV mini                                                 | 108-101  | mini/ spin/ vacuum | 100    |
|                                                                | 108-152  |                    | 250    |
| Clinic SV MAXI**                                               | 108-310  | MAXI/ spin/ vacuum | 10     |
|                                                                | 108-326  |                    | 26     |
| Plant SV mini                                                  | 117-101  | mini/ spin/ vacuum | 100    |
|                                                                | 117-152  |                    | 250    |
| Plant SV Midi**                                                | 117-226  | Midi/ spin/ vacuum | 26     |
|                                                                | 117-201  |                    | 100    |
| Plant SV MAXI**                                                | 117-310  | MAXI/ spin/ vacuum | 10     |
|                                                                | 117-326  |                    | 26     |
| GMO SV mini                                                    | 107-150  | mini/ spin/ vacuum | 50     |
|                                                                | 107-102  |                    | 200    |
| GMO SV MAXI**                                                  | 107-310  | MAXI/ spin/ vacuum | 10     |
|                                                                | 107-326  |                    | 26     |
| <b>GeneAll® GenEx™</b> <i>for the isolation of total DNA</i>   |          |                    |        |
| Genomic Sx <sup>†</sup>                                        | 208-001  | mini/ solution     | 100    |
| Genomic Sx <sup>†</sup>                                        | 208-005  | mini/ solution     | 500    |
| Genomic Lx <sup>††</sup>                                       | 208-301  | MAXI/ solution     | 100    |
| <b>GeneAll® RiboEx™</b> <i>for the isolation of total RNA</i>  |          |                    |        |
| RiboEx™                                                        | 301-001  | solution           | 100 ml |
|                                                                | 301-002  |                    | 200 ml |

\* GeneAll® Tissue SV mini, Midi, and MAXI plus! kit provide the additional methods for the purification from animal whole blood.

\*\* GeneAll® SV Midi/MAXI kits require the centrifuge which has a swinging-bucket rotor and ability of 4,000~5,000 xg.

<sup>†</sup> On the basis of DNA purification from 300 ul whole blood.

<sup>††</sup> On the basis of DNA purification from 10 ml whole blood.

# GeneAll® Exgene™ Tissue

## Brief mini protocol for an animal tissue

*\* Before use this protocol, we strongly recommend you first read carefully the detailed protocol at page 11*

### Disruption and lysis

1. Disrupt up to **25 mg** of tissue sample in **200 ul** of buffer **TL** by homogenization, grinding or mincing.
2. Add **20 ul** of **Proteinase K** and mix well.
3. Incubate at **56°C** for complete lysis.
4. (Optional) For RNA-free DNA, treat 4 ul of RNase A (100 mg/ml) and incubate for 2 min at RT.

### Binding

5. Add **400 ul** of buffer **TB** and mix well.
6. Apply the mixture into SV column.
7. Centrifuge for a min and replace the collection tube with new one.

### Washing

8. Apply **600 ul** of buffer **BW** into SV column.
9. Centrifuge for 30 sec and replace the collection tube with new one.
10. Apply **700 ul** of buffer **TW** into SV column.
11. Centrifuge for 30 sec, discard the filtrate and re-insert the collection tube back.
12. Centrifuge for 1 min at full speed.

### Elution

13. Place the SV column into a fresh 1.5 ml microcentrifuge tube.
14. Add **200 ul** of buffer **AE**, incubate for 2 min at RT and centrifuge for 1 min at full speed.



# GeneAll<sup>®</sup> Exgene<sup>™</sup> Tissue

## Brief Midi (MAXI) protocol for an animal tissue

*\* Before use this protocol, we strongly recommend you first read carefully the detailed protocol at page 22 or 29*

*\* The number in the parenthesis denote the amounts for MAXI kit*



### Disruption and lysis

1. Disrupt up to 100 (250) mg of tissue sample in 1 (3) ml of buffer TL by homogenization, grinding or mincing.
2. Add 100 (250)  $\mu$ l of Proteinase K and mix well.
3. Incubate at 64 °C for complete lysis.
4. (Optional) For RNA-free DNA, treat 15 (40)  $\mu$ l of RNase A (100 mg/ml) and incubate for 3 min at RT.



### Binding

5. Add 2 (7) ml of buffer TB and mix well.
6. Apply the mixture into SV column.
7. Centrifuge for 2 min and discard the filtrate.



### Washing

8. Apply 3 (7) ml of buffer BW into SV column.
9. Centrifuge for 2 min and discard the filtrate.
10. Apply 4 (10) ml of buffer TW into SV column.
11. Centrifuge for 15 min at full speed.



### Elution

12. Place the SV column into a fresh 15 (50) ml conical tube.
13. Add 300 (600)  $\mu$ l of buffer AE, incubate for 2 min at RT and centrifuge for 5 min at full speed.



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