

**FavorPrep™**  
**Plant Genomic**  
**DNA Extraction Mini Kit**

**User Manual**

**Cat. No.: FAPGK 001 (50 Preps)**  
**FAPGK 001-1 (100 Preps)**

**For Research Use Only**

## Introduction

Genomic DNA Mini Kit provides a fast and simple method to isolate total DNA (genomic DNA, mitochondrial and chloroplast) from plant tissue and cells. In the process, sample is disrupted by grinding in liquid nitrogen and lysis buffer incubation. The Lysate is treated with RNase A to degrade RNA and filtrated by filter column to remove cell debris and salt precipitations. In the presence of binding buffer with chaotropic salt, the genomic DNA in the lysate binds to glass fiber matrix in the spin column. The contaminants are washed with an ethanol contained wash buffer and finally, the purified genomic DNA is eluted by low salt elution buffer or water. The protocol does not require DNA phenol extraction and alcohol precipitation. The entire procedure can be completed in 60 minutes. The purified genomic DNA is ready for PCR, real-time PCR, Southern blotting and RFLP.

## Quality Control

The quality of Plant Genomic DNA Mini Kit is tested on a lot-to-lot basis. The Kits are tested by isolation of genomic DNA from 50 mg young leave. More than 10 µg of genomic DNA could be quantified with spectrophotometer and checked by agarose gel.

**Sample:** 100 mg of plant tissue

**Yield:** 5~30g

**Operation time:** <60 min

## Kit Contents

|                     | <b>FAPGK 001<br/>(50 preps)</b> | <b>FAPGK 001-1<br/>(100 preps)</b> |
|---------------------|---------------------------------|------------------------------------|
| FAPG1 Buffer        | 24ml                            | 50ml                               |
| FAPG2 Buffer        | 8ml                             | 15ml                               |
| FAPG3 Buffer*       | 15ml                            | 30ml                               |
| W1 Buffer**         | 22ml                            | 44ml                               |
| Wash Buffer***      | 10ml                            | 20ml                               |
| Elution Buffer      | 15ml                            | 30ml                               |
| RNase A (50mg/ml)   | 500 µl                          | 840 µl                             |
| Filter Column       | 50 pcs                          | 100 pcs                            |
| FAPG Column         | 50 pcs                          | 100 pcs                            |
| 1.5 ml Elution tube | 50 pcs                          | 100 pcs                            |
| 2ml Collection tube | 100 pcs                         | 200 pcs                            |

\*Add 30 ml/60 ml ethanol to FAPG3 Buffer before first use.

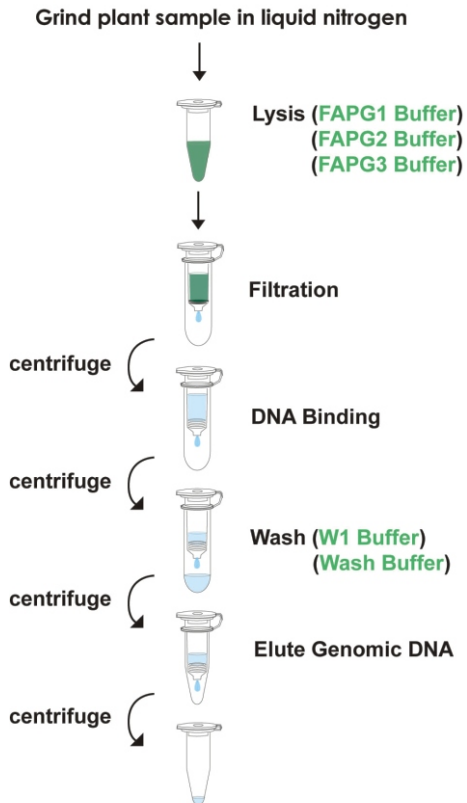
\*\*Add 8 ml/16ml ethanol to W1 Buffer before first use.

\*\*\*Add 40 ml/80ml ethanol to Wash Buffer before first use.

## Caution

The component contains irritant agent. During operation, always wear a lab coat, disposable gloves, and protective goggles.

# Procedure



## Protocol

|  |  |
|--|--|
| <p><b>Step 1<br/>Tissue<br/>Dissociation</b></p> | <ul style="list-style-type: none"> <li>• Cut off 50mg (up to 100mg) of fresh or frozen plant tissue or 5 mg (up to 100 mg) of dried sample.</li> <li>• Grind the sample under liquid nitrogen to a fine powder. Transfer it into a microcentrifuge tube (not provided). For some plant sample, we can destruct it without liquid nitrogen.</li> </ul>  |
| <p><b>Step 2<br/>Lysis</b></p>                   | <ul style="list-style-type: none"> <li>• Add 400µl <b>FAPG1 Buffer</b> and 4 µl <b>RNase A (100 mg/ ml)</b> into the sample tube and mix by vortexing. Do not mix FAPG1 Buffer and RNase A before use.</li> <li>• Incubate at 65°C for 10 minutes. During incubation, invert the tube every 5 minutes. At the same time, preheat required Elution Buffer (200 µl per sample) at 65°C.</li> <li>• Add 130µl <b>FAPG2 Buffer</b> and mix by vortexing.</li> <li>• Incubate at ice for 5 minutes.</li> <li>• Place a <b>Filter Column</b> in a 2 ml Collection Tube.</li> <li>• Apply the mixture from previous step to the <b>Filter Column</b>. Centrifuge for 3 minutes at full speed (13,000 rpm).</li> <li>• Discard the <b>Filter Column</b> and carefully transfer clarified supernatant in Collection Tube to a new microcentrifuge tube (not provided).</li> </ul> |
| <p><b>Step 3<br/>DNA Binding</b></p>             | <ul style="list-style-type: none"> <li>• Add 1.5 volumes of <b>FAPG3 Buffer (ethanol added)</b> to the cleared lysate and mix immediately by vortexing for 5 seconds. For example, add 750µl FAPG3 Buffer to 500 l lysate.</li> <li>• Place a <b>FAPG Column</b> in a 2 ml Collection Tube.</li> <li>• Apply 700µl the mixture (including any precipitate) from previous step to the FAPG Column.</li> <li>• Centrifuge at full speed (13,000 rpm) for 2 minute.</li> <li>• Discard flow-through in Collection Tube and apply remaining mixture to FAPG Column.</li> <li>• Centrifuge at full speed (13,000 rpm) for 2 minute.</li> <li>• Discard flow-through in Collection Tube.</li> </ul>  |

|                                      |  |
|--------------------------------------|--|
| <p><b>Step 4<br/>Wash</b></p>        | <ul style="list-style-type: none"> <li>• Add 500µl of <b>W1 Buffer (ethanol added)</b> into the column.</li> <li>• Add 750µl of <b>Wash Buffer (ethanol added)</b> into the column.</li> <li>• Centrifuge at full speed (13,000 rpm) for 30 seconds.</li> <li>• Discard the flow-through and place the FAPG Column back in the Collection Tube.</li> <li>• Discard the flow-through and place the FAPG Column back in the Collection Tube.</li> <li>• Centrifuge at full speed for 3 minutes to dry the column matrix.</li> </ul>  |
| <p><b>Step 5<br/>DNA Elution</b></p> | <ul style="list-style-type: none"> <li>• Standard elution volume is 100µl. If less sample to be used, reduce the elution volume (30-50µl) to increase DNA concentration. If higher DNA yield is required, repeat the DNA Elution step to increase DNA recovery and the total elution volume is about 200µl.</li> <li>• Transfer dried <b>FAPG Column</b> into a clean 1.5 ml microcentrifuge tube (not provided).</li> <li>• Add 100µl of <b>preheated Elution Buffer</b> into the center of the column matrix.</li> <li>• Stand for 3-5 minutes until Elution Buffer absorbed by the matrix.</li> </ul> <p>Centrifuge full speed (13,000 rpm) for 30 seconds to elute purified DNA.</p> |

# Troubleshooting

| Problem                       | Possible Reasons/ Solution  |
|-------------------------------|---|
| <p><b>Low yield</b></p>       | <p><b>Insufficient Lysis</b></p> <ul style="list-style-type: none"> <li>• Prolong the incubation time in lysis buffer to obtain higher yields of DNA.</li> </ul>  |
|                               | <p><b>Insufficient disruption</b></p> <ul style="list-style-type: none"> <li>• For most of species we recommend grinding with liquid nitrogen. Homogenization should be done thoroughly until the plant material is ground to a fine powder.</li> </ul>   |
|                               | <p><b>DNA still bound to the membrane</b></p> <ul style="list-style-type: none"> <li>• The DNA can be either eluted in higher volumes or by repeating the elution step up to three times. Elution buffer should be preheated to 60°C prior to elution.</li> <li>• To ensure correct pH, use supplied elution buffer.</li> </ul> |
| <p><b>DNA is degraded</b></p> | <p><b>Sample was contaminated with DNase</b></p> <ul style="list-style-type: none"> <li>• Preheat elution buffer to 60°C for 5 minutes to eliminate any possible Dnase</li> </ul>   |
|                               | <p><b>Centrifugation speed was too high</b></p> <ul style="list-style-type: none"> <li>• Higher velocities may cause shearing of the DNA. The centrifugation maximum speed is at 11,000xg.</li> </ul>   |
| <p><b>Column clogged</b></p>  | <p><b>Too much tissue was used.</b></p> <ul style="list-style-type: none"> <li>• Too much tissue was used. Reduce the amount of sample material or separate it into multiple tubes.</li> </ul>  |
|                               | <p><b>Insufficient centrifugation</b></p> <ul style="list-style-type: none"> <li>• Centrifuge again and extend centrifugation time.</li> </ul>  |
|                               | <p><b>Precipitate was formed at DNA Binding Step</b></p> <ul style="list-style-type: none"> <li>• Reduce the sample material.</li> <li>• Before loading the column, break up the precipitate in ethanol-added lysate by pipetting.</li> </ul>   |