

# FavorPrep™ Blood Genomic DNA Extraction Mini Kit

Cat.No.: FABG1020 (4 preps) FABG1023 (50 preps) FABG1024 (100 preps) FABG1026 (300 preps) (For Research Use Only)

Kit Contents:	FABG1020 (4 preps)	FABG1023 (50 preps)	FABG1024 (100 preps)	FABG1026 (300 preps)
FABG Buffer	1.5 ml	15 ml	30 ml	70 ml
W1 Buffer * (Concentrate)	1.3 ml	22 ml	44 ml	124 ml
Wash Buffer ** (Concentrate)	1 ml	10 ml	20 ml	50 ml
Elution Buffer	1 ml	15 ml	30 ml	90 ml
Proteinase K (Liquid)	100 μΙ	1050 µl	1050 µl × 2	1600 µl × 4
FABG Mini Columns	4 pcs	50 pcs	100 pcs	300 pcs
Collection Tubes	8 pcs	100 pcs	200 pcs	600 pcs
Elution Tubes	4 pcs	50 pcs	100 pcs	300 pcs
User Manual	1	1	1	1

* Preparation of W1 Buffer and Wash Buffer for the first use:					
Cat. No:	FABG1020 (4 preps)	FABG1023 (50 preps)	FABG1024 (100 preps)	FABG1026 (300 preps)	
Ethanol volume for W1 Buffer *	0.5 ml	8 ml	16 ml	45 ml	
Ethanol volume for Wash Buffer **	4 ml	40 ml	80 ml	200 ml	

# Specification:

Principle: spin column (silica membrane)

Sample: up to 200 µl whole blood, serum, plasma, body fluids

up to 5×106 cultured cells

Operation time: <30 mins

Binding capacity: up to 60  $\mu$ g/column DNA Yield: 4~8  $\mu$ g/200  $\mu$ l whole blood

Elution Volume: 50~200 ul

### **Important Notes:**

- 1. Buffers provided in this system contain irritants. Wear gloves and lab coat when handling these buffers.
- 2. Add required volume of ethanol (96~100%) to W1 Buffer and Wash Buffer at the first use.
- 3. Preheat a dry bath or water bath to 60°C and 65°C before the operation.
- 4. All centrifuge steps are done at full speed (~18,000 xg) in a microcentrifuge.

# Sample optional: Add RNase A that RT, 2 mins Cell lysis (FABG Buffer) (Proteinase K) that 60°C, 15 mins Add ethanol Centrifuge Wash (W1 Buffer) (Wash Buffer) Centrifuge Furified genomic DNA

## **General Protocol:**

Hints: Prepare a dry bath or water bath to 60°C bath for step 4. Preheat Elution Buffer to 65°C for step 13 (Elution step).

### Please Read Important Notes Before Starting the Following Steps.

- 1. Transfer up to 200 µl sample (whole blood, serum, plasma, body fluids, buffy coat) to a microcentrifuge tube (not provided).

  -If the sample volume is less than 200 µl, add the appropriate volume of PBS.
- 2. **(Optional)**: If RNA-free genomic DNA is required, add 4 µl of 100 mg/ml RNase A to the sample and incubate for 2 mins at room temperature.
- 3. Add 20 µl Proteinase K and 200 µl FABG Buffer to the sample. **Mix thoroughly by pulse-vortexing.**-Do not add Proteinase K directly to FABG Buffer.
- 4. Incubate at 60°C for 15 mins to lyse the sample. During incubation, vortex the sample every 3~5 mins interval.
- 5. Briefly spin the tube to remove drops which inside of the lid.
- 6. Add 200 µl ethanol (96~100%) to the sample. Mix thoroughly by pulse-vortexing for 10 secs.
- 7. Briefly spin the tube to remove drops which inside of the lid.
- 8. Place a FABG Mini Column to a Collection Tube. Transfer the mixture (including any precipitate) carefully to the FABG Mini Column. Centrifuge at 6,000 xg for 1 min then place FABG Mini Column to a new Collection Tube.
- 9. Add 400 µl W1 Buffer to the FABG Mini Column and centrifuge at full speed for 30 secs then discard the flow-through.

  -Make sure that ethanol has been added into W1 Buffer at the first open.
- 10. Add 750 µl Wash Buffer to the FABG Mini Column and centrifuge at full speed for 30 secs then discard the flow-through.

  -Make sure that ethanol has been added into Wash Buffer at the first open.
- 11. Centrifuge at full speed for an additional 3 mins to dry the column.
  - Important Step! This step will avoid the residual liquid to inhibit subsequent enzymatic reaction.
- 12. Place the FABG Mini Column to an Elution Tube.
- 13. Add 50~200 µl of heated Elution Buffer or ddH2O (pH 7.5~9.0) to the membrane center of FABG Mini Column. **Stand FABG Mini Column for 3 mins.** 
  - -Important Step! For effective elution, make sure that the elution solution is dispensed onto the membrane center and is absorbed completely.
- 14. Centrifuge at full speed for 1 min to elute total DNA.
- 15. Store total DNA at 4°C or -20°C.

# **Special Protocol:**

### For cultured cells

- 1. Harvest cells
  - a. Cells grown in suspension
    - i. Transfer the appropriate number of cells (up to 5×106) to a 1.5 ml microcentrifuge tube.
    - ii. Centrifuge at 300 x g for 5 mins.
  - iii. Remove the supernatant carefully and completely.
  - b. Cells grown in monolayer
    - i. Detach cells from the dish or flask by trypsinization or using a cell scraper.
    - ii. Transfer the appropriate number of cells (up to 5×10°) to a 1.5 ml microcentrifuge tube.
    - iii. Centrifuge at 300 xg for 5 mins.
    - iv. Remove the supernatant carefully and completely.
- 2. Resuspend cell pellet in PBS to a final volume of 200 µl.
- 3. Follow the General Protocol starting from step 2.

# Preparation of buffy coat

Centrifuge whole blood at 3,300 xg for 10 mins at room temperature and you will get three different fractions: the upper clear layer is plasma; the intermediate layer is buffy coat, containing concentrated leukocytes; the bottom layer contains concentrated erythrocytes. Process the General Protocol from Step 1 for buffy coat. Extraction total DNA from buffy coat will yield 5~10 times more DNA than an equivalent volume of whole blood.

# **Troubleshooting**

Possible reasons	Solutions				
Low or no yield of genomic DNA					
Low amount of cells in the sample	Concentrate a larger volume of a new sample to 200 µl. If the sample is whole blood, prepare buffy coat.				
Poor cell lysis					
Poor cell lysis because of insufficient Proteinase K activity	Repeat the extraction procedure with a new sample. Make sure the reactive temperature and time is correct.				
Poor cell lysis because of insufficient mixing with FABG buffer	Repeat the extraction procedure with a new sample. Mix the sample and FABG Buffer immediately and thoroughly by pulse-vortexing.				
Poor cell lysis because of insufficient incubation time	Repeat the extraction procedure with a new sample. Extend the incubation time and make sure that no residual particulates remain.				
Ethanol is not added into the lysate before transferring into FABG Mini Column	Repeat the extraction procedure with a new sample.				
Incorrect preparation of Wash Buffer					
Ethanol is not added into Wash Buffer at the first open	Make sure that the correct volume of ethanol (96~100%) is added into V				
The volume or the percentage of ethanol is not correct before adding into Wash Buffer	Buffer at the first open. Repeat the extraction procedure with a new sample.				
Elution of genomic DNA is not efficient					
pH of water (ddH2O) for elution is acidic	Make sure the pH of ddH2O is between 7.5~9.0.				
	Use Elution Buffer (provided) for elution.				
Elution Buffer or ddH <sub>2</sub> O is not completely absorbed by column membrane	After Elution Buffer or ddH <sub>2</sub> O is added, stand the FABG Mini Column for 5 mins before centrifugation.				
Column is clogged					
Blood sample contains clots	Repeat the extraction procedure with a new sample. Mix the blood samp well with anti-coagulant to prevent formation of blood clots.				
Sample is too viscous	Reduce the sample volume.				
Degradation of elutated DNA					
Sample is old	Always use fresh or well-conserved sample for genomic DNA extraction.				
Buffer for gel electrophoresis contaminated with DNase	Use fresh running buffer for gel electrophoresis.				