

N[®] FavorPrep[™] Blood/Cultured Cell Genomic DNA Extraction Maxi Kit

(For Research Use Only)

Kit Contents:

Cat. No. (preps)	FABGK000-Maxi (2 preps)	FABGK003 (10 preps)	FABGK003-1 (24 preps)
Proteinase K (Liquid)	1050 µl × 2	10.5 ml	13 ml × 2
FABG Buffer	22 ml	110 ml	265 ml
W1 Buffer* (Concentrate)	6.5 ml	33 ml	88 ml
Wash Buffer** (Concentrate)	3 ml	20 ml	40 ml
Elution Buffer	6 ml	30 ml	60 ml
FABG Maxi Column	2 pcs	10 pcs	24 pcs
Elution Tube (50 ml tube)	2 pcs	10 pcs	24 pcs
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Preparation of W1 Buffer and Wash Buffer for the first use:

Cat. No:	FABGK000-Maxi (2 preps)	FABGK003 (10 preps)	FABGK003-1 (24 preps)
* Ethanol volume for W1 Buffer	2.5 ml	12 ml	32 ml
**Ethanol volume for Wash Buffer	12 ml	80 ml	160 ml

Specification:

Principle: spin column (silica membrane) Sample Size: up to 10 ml of fresh/frozen blood up to 1×10⁸ of cultured cells Column Capacity: 500 µg of DNA Average DNA yield: 35 µg/ml of whole blood Handling Time: 1 hr Elution Volume: 0.75~1.5 ml

Required material to be provided by user

Pipettors and pipet tips Centrifuge: should be capable up to 4,000 x g Thermal incubator Oven (optional) Ethanol (96~100%) Vortex



Important Notes:

- 1. Buffers provided in this system contain irritants. Wear gloves and lab coat when handling these buffers.
- 2. Preheat a thermal incubator to 60°C before the operation.
- 3. Use a centrifuge with a swinging bucket rotor and a force of 4,000~6,000 x g for in all centrifugation steps.
- 4. Preheat the Elution Buffer or ddH2O for step 11 (Elution step).

Protocol: For Blood DNA Extraction

Please Read Important Notes Before Starting the Following Steps.

- 1. Transfer up to 10 ml sample (whole blood, buffy coat) to a 50 ml centrifuge tube (not provided). If the sample volume is less than 10 ml, add PBS to adjust volume to 10 ml.
- Add 1000 µl of Proteinase K to the sample and mix well by vortexing. Add 10 ml of FABG Buffer to the sample mixture. Mix thoroughly by pulse-vortexing.
 Do not add Proteinase K directly to FABG Buffer.
- Incubate the sample mixture at 60°C for 15 mins to lyse the sample. During incubation, invert the tube every 3~5 mins.
- 4. **(Optional)**: If RNA-free genomic DNA is required, add 80 µl of 100 mg/ml RNase A (not provided) to the sample mixture and incubate at room temperature for 10 mins.
- 5. Add 10 ml of ethanol (96~100%) to the sample mixture. Mix thoroughly by vortexing; If precipitate appears, break it by pipetting.
- 6. Place a FABG Maxi Column to a 50 ml centrifuge tube (not provided). Transfer 15.5 ml of sample mixture (ethanol added; Including any precipitate) carefully to the FABG Maxi Column. Close the cap and **centrifuge** at 4,000~6,000 x g for 3 mins.

- 7. Discard the flow-through and transfer the rest sample mixture to the same FABG Maxi Column. Close the cap and **centrifuge at 4,000~6,000 x g for 3 mins** and discard the flow-through.
- 8 Add 4 ml of W1 Buffer (ethanol added) to the FABG Maxi Column. Close the cap and **centrifuge at 4,000~6,000 x g for 3 mins**. Discard the flow-through and place the FABG Maxi Column back in the 50 ml centrifuge tube. -Make sure that ethanol has been added into W1 Buffer at the first open.
- Add 7 ml of Wash Buffer (ethanol added) to the FABG Maxi Column. Close the cap and centrifuge at 4,000~
 6,000 x g for 15 mins. Discard the flow-through and place the FABG Maxi Column back in the 50 ml centrifuge tube.

-Make sure that ethanol has been added into Wash Buffer at the first open.

-Important Step! Make sure the residual liquid have been removed completely after centrifugation. It might be necessary to do a further drying by placing the column in a vacuum oven at 70°C for 3 mins.

- 10. Place the FABG Maxi Column into a new 50 ml centrifuge tube (Elution Tube, provided).
- 11. Add 0.75~1.5 ml of preheat Elution Buffer or ddH₂O (pH 7.5-9.0) to the membrane center of the FABG Maxi Column. Stand the FABG Maxi Column for 5 mins at room temperature.

-Important Step! For effective elution, stand the FABG Maxi Column for 5 mins is required to make sure Elution Buffer is absorbed completely by column membrane.

12. Centrifuge at 4,000 x g for 2 mins to elute total DNA. -Standard volume for elution is 1 ml. If higher DNA yield is required, repeat the DNA Elution step (step 11 and

12) to increase DNA recovery.

Protocol: For Cultured Cell DNA Extraction

- 1. Transfer up to 1×10⁸ of cells to a 50 ml centrifuge tube (not provided). **Centrifuge at 4,000~6,000 x g for 5 mins** to pellet the cells. (If using adherent cells, trypsinize the cells before harvesting.)
- 2. Resuspend the cells with 10 ml of PBS.
- 3. Follow the Blood protocol starting from step 2.

Troubleshooting

Possible reasons	Solutions	Possible reasons	Solutions	
Low or no yield of genomic DNA		Elution of genomic DNA is not efficient		
Poor cell lysis		pH of water (ddH2O) for	Make sure the pH of ddH2O is between 7.5-9.0.	
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		Use Elution Buffer (provided) for elution.	
Poor cell lysis because of insufficient mixing with FABG buffer	cell lysis because of ficient mixing with Repeat the extraction proce-dure with a new sample. Mix the sample and FABG Buffer G buffer sample and FABG Buffer		After Elution Buffer or ddH2O is added, stand the FABG Maxi Column for 5 mins before centrifugation.	
	pulse-vortexing.	Column is clogged		
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.	Blood sample contains clots	Repeat the extraction proce- dure with a new sample. Mix the blood sample well with anti-co- agulant to prevent formation of blood clots.	
Ethanol is not added into	Repeat the extraction procedure	Sample is too viscous	Reduce the sample volume.	
the lysate before transfer-	with a new sample.	Degradation of elutated DNA		
Column	b Buffor	Sample is old	Always use fresh or well-con- served sample for genomic DNA	
Ethemal is not asked into the table sure that the sourcest			extraction.	
W1 and Wash Buffer at the first open	volume of ethanol (96~100%) is added into W1 and Wash Buffer at the first open. Repeat the	Buffer for gel electropho- resis contaminated with DNase	Use fresh running buffer for gel electrophoresis.	
	extraction procedure with a new sample.			
The volume or the percentage of ethanol is not correct before adding into W1 and Wash Buffer	Make sure that the correct volume of ethanol (96~100%) is added into W1 and Wash Buffer at the first open. Repeat the extraction procedure with a new sample.			