

# FavorPrep™ Endotoxin-Free Plasmid Extraction Maxi Kit

For Research Use Only

| Kit Contents:    | FAPD2050<br>(2 preps) | FAPD2051<br>(10 preps) |
|------------------|-----------------------|------------------------|
| PEQ Buffer       | 30 ml                 | 135 ml                 |
| PM1 Buffer       | 42 ml                 | 215 ml                 |
| PM2 Buffer       | 42 ml                 | 215 ml                 |
| PM3 Buffer       | 42 ml                 | 215 ml                 |
| PTR Buffer       | 12 ml                 | 55 ml                  |
| PW Buffer        | 65 ml                 | 270 ml + 60 ml         |
| PEL Buffer       | 32 ml                 | 215 ml                 |
| RNase A Solution | 180 µl                | 900 µl                 |
| PM Maxi Columns  | 2 pcs                 | 10 pcs                 |

| * Preparation of PM1 Buffer for the first use: |                       |                        |  |
|--|-----------------------|------------------------|--|
| Cat. No:                                       | FAPD2050<br>(2 preps) | FAPD2051<br>(10 preps) |  |
| Volume of RNase A<br>Solution for PM1 Buffer   | 168 µl                | 860 µl                 |  |

# **Specification:**

Technology: Anion-exchange chromatography (gravity-flow column)

Lysate clarification: Centrifugation

Sample Size: 120~240 ml of bacteria for high-copy number or low

-copy number plasmid

Plasmid or constructs range: 3 kbp~150 kbp Binding Capacity: 1.5 mg/Maxi Column

## **Important Notes:**

- 1. Store RNase A Solution at -20°C upon receipt of kit.
- Add indicated volume of RNase A Solution into PM1 buffer, mix well and store the PM1 buffer at 4°C.
- 3. If precipitates have formed in PM2 Buffer, warm the buffer in 37°C water bath to dissolve precipitates.
- 4. Pre-chill PM3 Buffer at 4°C before starting.

# **Additional Requirements:**

- 1. 50 ml tubes
- 2. Refrigerated centrifuge capable of ≥5,000 xg and the centrifuge tube suitable for the centrifuge rotor
- 3. Isopropanol
- 4. 70% ethanol
- 5. TE buffer or ddH2O

# **General Protocol:**

Please Read Important Notes Before Starting Following Steps.

### Harvest bacterial cells

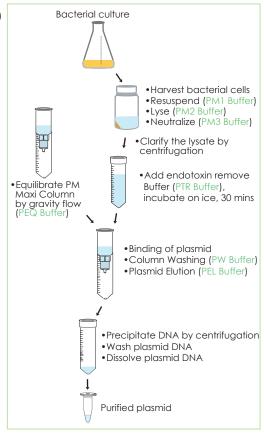
1. Harvest the cells by centrifugation at 4,500~6,000 xg at 4°C for 10 mins and discard the supernatant.

### Equilibrate PM Maxi Column

- 2. Place a PM Maxi Column onto a 50 ml tube.
- 3. Equilibrate the PM Maxi column by applying 10 ml of PEQ Buffer. Allow the column to empty by gravity flow and discard the filtrate.

## Cell lysis and lysate neutralization

- 4. Add 16 ml of PM1 Buffer (RNase A added) to resuspend the cell pellet by vortexing or pipetting.
- 5. Add 16 ml of PM2 Buffer and mix gently by inverting the tube 5 times.
  - -Do not vortex to avoid shearing genomic DNA.
- 6. Incubate the sample mixture for 5 mins at room temperature until lysate clears.



- 7. Add 16 ml of chilled PM3 Buffer and mix immediately by inverting the tube 10~15 times to neutralize the lysate. (Do not vortex!)
  - -Note: •Make sure the density of cultured cell is optimal, the buffers volume (PM1, PM2, PM3) should be increased proportionally to the culture volume.
    - (ex. culture volume, 120~240 ml: PM1, 16 ml; PM2, 16 ml; PM3, 16 ml culture volume, 240~480 ml: PM1, 32 ml; PM2, 32 ml; PM3, 32 ml)
    - When processing 240~480 ml bacteria where additional buffer volumes of PM1, PM2, and PM3 are needed, the buffers may be purchased separately.
    - Make sure cell pellet be suspended completely within Buffer PM1.
    - •Mix the sample mixture completely after adding Buffer PM2 and Buffer PM3.

## Lysate clarification and endotoxin removal

- 8. Centrifuge the tube at ≥5,000 xg at 4°C for 20 mins. (Preferably centrifuge the tube at 15,000~20,000 xg at 4°C for 15 mins.)
  - -If the supernatant still contains suspended matter, transfer the supernatant to a clean centrifuge tube and repeat this centrifugation step.
- 9. Transfer the supernatant to a clean 50 ml tube.
- 10. Add 5 ml of PTR Buffer and mix gently by pipetting. Incubate the sample mixture on ice for 30 mins. After the incubation, the sample mixture will become clear.

## Binding of plasmid

- 11. Transfer the half of the sample mixture from step 10 to the equilibrated PM Maxi Column. Allow sample mixture to flow through the PM Maxi Column by gravity flow and discard the filtrate.
- 12. Repeat step 11 for the rest of the sample mixture.

### Wash PM Maxi Column

13. Wash the PM Maxi column by applying 30 ml of PW Buffer. Allow PW Buffer to flow through the PM Maxi Column by gravity flow and discard the filtrate.

#### **Elution**

14. Place the PM Maxi column onto a clean 50 ml centrifuge tube (not provided). Add 15 ml of PEL Buffer to the PM Maxi Column to elute the plasmid by gravity flow.

### Precipitate plasmid DNA

- 15. Transfer the eluate from step 14 to a centrifuge tube. Add 0.75X volume of room temperature isopropanol to the eluate and mix well by inverting the tube 10 times. (ex: add 11.25 ml isopropanol to 15 ml eluate)

  -Note! Make sure that isopropanol be mixed thoroughly with eluate before centrifugation.
- 16. Centrifuge the tube at ≥5,000 xg at 4°C for 30 mins. (Preferably centrifuge the tube at 15,000~20,000 xg at 4°C for 20 mins.)

### Wash and dissolve plasmid DNA

- 17. Carefully remove the supernatant and wash the plasmid pellet with 5 ml of room temperature 70% ethanol.
- 18. Centrifuge the tube at ≥5,000 xg at 4°C for 10 mins.
- 19. Carefully remove the supernatant and invert the tube on paper towel for 3 mins to remove residual ethanol. Air-dry the plasmid pellet until the tube is completely dry. (Or incubate the plasmid pellet at 70°C for 10 mins.)
- 20. Dissolve the plasmid pellet in a suitable volume (≥300 µl) of TE or ddH2O.
  - -Note! Do not lose the DNA pellet when discard the supernatant.
    - Make sure the DNA pellet adhesive lightly on the centrifuge tube.
    - If the DNA pellet loose from tube, repeat the precipitation step again.
    - Make sure the DNA is dissolved completely before measure the concentration.

# **Troubleshooting**

### Low yield

## Bacterial cells were not lysed completely

- •Too many bacterial cells were used.
- After PM3 Buffer addition, break up the precipitate by inverting.
- DNA failed to precipitate or DNA pellet was lost after precipitation.
- DNA pellet was insufficiently redissolved.

# Purified DNA dose not perform well in downstream applications

### RNA contamination

- Make sure that RNase A had been added in PM1 Buffer at the first use.
- RNase A is not properly preserved.
- •Too many bacterial cells were used, reduce the sample volume.

# Genomic DNA contamination

- •Do not use overgrown bacterial culture.
- During PM2 and PM3 Buffer addition, mix gently to prevent genomic DNA shearing.
- •Lysis time was too long (over 5 mins).

## Too much salt residual in DNA pellet

•Wash the DNA pellet twice with 70% ethanol.