

ExiProgen™EC-Tagfree Protein Synthesis Kit (Cat. No. K-7320)

Step | Preparation of Template DNA

1) Prepare the 'Template DNA' for protein expression.

(Note I) The 'Template DNA' must contain the following: T7 Promoter, Ribosome binding site (RBS), His tag-TEV cleavage site - Target gene, and T7 terminator.

(Note II) Use 0.6 μg/kb of plasmid DNA (Purity: A260/280 >1.7, A260/230 >1.5).

Step | Preparation of Experiment

- 1) Take out the 'Cartridge 2' from Box 2. Thaw them at room temperature.
- 2) Take out 'E. coli extract,' Master Mix,' DEPC DW', 'Storage Buffer' and 'TEV protease' from **Box** ② and thaw them on ice.

 $\label{thm:continuous} {\sf Take\ out'Cartridge\ \textcircled{1},'Disposable\ Filter\ Tip,' and'Protection\ Cover' from\ \textbf{Box\ \textcircled{1}}.}$

 $\textbf{(Note)} \ \textit{Make sure that all solutions are completely thawed.} \ \textit{It takes about 2 hours.}$

- 3) Prepare the 'Reaction Block (ExiProgen™ accessory)'.
- 4) Take out twice as many 'Dialysis Tube' as the number of samples from **Box** 1.
- 5) Remove the solution in the 'Dialysis Tube' and rinse inside and out using squeeze bottle with sterile water.

(Note) The solution in the 'Dialysis Tube' is 20%(v/v) ethanol.

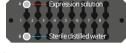
6) Install the 'Dialysis Tube' on the 'Reaction Block,' as shown in the diagram.



7) Remove the water in the 'Dialysis Tube' and prepare expression solution as shown in the table. The expression solution should be loaded at 'Dialysis Tube' in row A (Protein expression section). Fill 500 µl of sterile distilled water in the 'Dialysis Tube' on row B (Dialysis section).

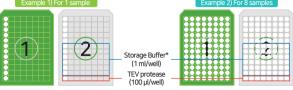
*Composition of Expression Solution

| | | Positive control DNA (Optional) | |
|----------------|------------|------------------------------------|--|
| Template DNA | Xμl | 6 µl | |
| E.coli extract | 120 µl | 120 µl | |
| Master Mix | 210 µl | 210 µl | |
| DEPC DW | (120-X) µl | 114 µl | |
| Total volume | 450 µl | 450 µl | |



- 8) Place the 'Protection Cover' on the 'Reaction Block'.
- 9) Punch holes in the sealing films of 'Cartridge 1), 2).
- 10) Add 1 $\underline{\text{ml}}$ of 'Storage Buffer' to G2-J1 rows and 100 μl of 'TEV protease' to G1 row of 'Cartridge 2).

(Note) The number of columns punched should match the number of the samples.



* Composition of Storage Buffer

 $-50\,\mathrm{mM\,Tris}\text{-HCl,}\,100\,\mathrm{mM\,NaCl,}\,1\,\mathrm{mM\,DTT,}\,0.1\,\mathrm{mM\,EDTA},\\ 0.05\%(v/v)\,\mathrm{NaN3,}\,50\%(v/v)\,\mathrm{glycerol,}\,\mathrm{pH}\,7.6\,\mathrm{mM\,Tris}$

Step III Protein Synthesis with ExiProgen™

Turn on the ExiProgen™ and tap 'Press to Start' button.
Wait until the [MENU] screen appears.

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- 2) Open the door of ExiProgen™ and pull out the baseplate.
- 3) Load 'Cartridge 1), 2), 'Waste Tray', and 'Reaction Block'.

Follow in order: 'Cartridge ②' → 'Cartridge ①' → 'Waste Tray' → 'Reaction Block'.

(Note I) Make sure that the first and second lines of 'Cartridge @' are firmly fixed on the heating block.

(Note II) There are 'silicon rings' embedded on both sides of the 'Cartridge ①' installation position. Place the left side first, and then the right side.

4) Place the 'Disposable Filter Tips' on row B of 'Disposable Tip Rack'.

(Note) Tips should be placed at the same columns with the punched holes of the Cartridges.













- 5) Push the baseplate in until you hear the click sound, then close the door.
- 6) Tap the following: [MENU] 'Start' → [PREP SETUP] '904 (Synthesis_TF), 'Enter' → [PREP SETUP, Elution Volume] 'OK' → [PREP SETUP, Reaction Temperature] '30°C, 'OK' → [CHECK LIST] 'OK' → [Running Mode] 'RUN'.













- The [Work Completion] screen will appear once the protocol is completed. It takes approximately 26 hours.
- 8) Purified protein samples in the 'Storage Buffer' can be collected from the 'Dialysis Tube', located at the row B of the 'Reaction Block'.

Step IV Analysis of Sample

A. Position of each sample (in 'Cartridge (2)')



IExpression sample

: Samples after expression, but without purification | Unbound sample

: Samples not bound to Ni-NTA magnetic bead IBead sample

: Used bead samples for purification

B. Sampling for SDS-PAGE analysis

1) Prepare the loading mixture as shown in the table. Incubate the samples at 95°C for 5-10 min.

| | Expression/ Unbound sample | Purified protein sample | Bead sample |
|-------------------------|-------------------------------|-------------------------|-------------|
| Sample | 5 µl | 5 µl | 15 µl |
| 4X Loading dye | 5 µl | 5 µl | 5 µl |
| Sterile distilled water | 10 µl | 10 µl | - |
| Total volume | 20 µl | 20 µl | 20 µl |

(Note) Add 360-400 μ l of sterile distilled water to the well containing the beads for suspension. Proceed with the sampling afterwards.

Load each sample to the wells of SDS-PAGE gel [10 x 8 (cm), 0.75 mm thick, 10-well].
(Note) Expression, Unbound sample: 5 µl/well,

Purified proteins and Bead samples: 10 µl/well For more information, visit our website (www.bioneer.com).

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