





# **GeneMATRIX** Basic DNA Purification Kit

3 in 1: For Agarose, Plasmid and PCR/DNA Clean-Up

Universal kit for purification of PCR products / DNA after enzymatic reactions, isolation of DNA from agarose gels and isolation of plasmid DNA from bacteria.

O Cat. no. E3545

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| Content             | 50 preps<br>E3545-01 | 150 preps<br>E3545-02 | Storage/Stability |  |  |  |
|---------------------|----------------------|-----------------------|-------------------|--|--|--|
| Buffer Uni          | 1.8 ml               | 5.4 ml                | 15-25°C           |  |  |  |
| Basic               | 60 ml                | 180 ml                | 15-25°C           |  |  |  |
| Cell R *            | 15 ml                | 45 ml                 | 2-8°C             |  |  |  |
| Lysis Blue          | 15 ml                | 45 ml                 | 15-25°C           |  |  |  |
| Neutral B           | 21 ml                | 63 ml                 | 15-25°C           |  |  |  |
| Wash UX1            | 30 ml                | 90 ml                 | 15-25°C           |  |  |  |
| Wash UX2            | 36 ml                | 108 ml                | 15-25°C           |  |  |  |
| Elution             | 9 ml                 | 27 ml                 | 15-25°C           |  |  |  |
| DNA Binding Columns | 50                   | 3 x 50                | 15-25°C           |  |  |  |
| Protocol            | 1                    | 1                     |                   |  |  |  |

<sup>\*</sup> Contains lysozyme (20 mg/ml).

# **Introductory Notes**

### **NOTE 1** · Kit Specification.

PCR/DNA Clean-up Protocol is suitable for fast cleanup of up to 25  $\mu$ g of DNA fragments from PCR and others enzymatic reactions (sizes from approximately 100 bp to over 15 kb). Procedure removes primers below 40 nt and double-stranded DNA below 20 bp. However, common short by-products of not optimal or problematic PCR, known as primer-dimers, also consist of double-stranded DNA. They are produced from self-annealed and extended primers and co-migrate on a gel along with unincorporated single-stranded DNA primers. These double-stranded DNA artefacts co-purify with an expected PCR product, if their length exceeds 20 bp. If the removal of primer-dimers is necessary, we recommend PCR reaction optimization and/or agarose gel electrophoresis followed by isolation of PCR product using Agarose-Out Protocol.

**Agarose-Out Protocol** is designed to isolate DNA molecules, ranging in size from approximately 100 bp to 10 kb, from TAE- or TBE-agarose gels. It is also possible to purify DNA fragments up to 20 kb or more, with decreased isolation yields. Up to 250 mg agarose can be processed per spin column.

**Plasmid DNA Purification Protocol** is designed to isolate high purity plasmid DNA from various species of Gram- bacteria, including recombinant Escherichia coli strains. Recommended culture volume is 1.0–3.0 ml. Use either stationary phase or log phase bacterial cultures for obtaining highest DNA quality. Due to differences in growth characteristics of bacteria species, it is recommended to perform a preliminary experiment for determining the optimal starting amount. In general, the weight of the cell pellet should not exceed 50 mg per single minicolumn and the volume of the culture volume should not exceed 3.0 ml per single minicolumn. In case of minicolumn clogging due to high lysate viscosity, reduce the initial amount of bacteria used for isolation.

**NOTE 2 · Maximum Sample Amount.** The maximum column binding capacity for DNA is 25 μg. The maximum volume of the column reservoir is 650 μl.

**NOTE 3** • **Kit Compounds Storage**. Once the kit is unpacked, store components at room temperature, except buffer Cell R, which should be stored at 2–8°C. This will ensure the best performance, due to preserving activity of RNaseA included into the buffer. In case of occasional buffer ingredients precipitation, simply warm up in 37°C water bath, until clarified.

**NOTE 4. Maintaining Good Working Practice.** All solutions should be kept tightly closed to avoid evaporation and resulting components concentration changes. Buffer Basic may form reactive and toxic compounds when combined with acids. Do not add bleach or acidic solutions to the sample preparation waste.

# Equipment and reagents to be supplied by user:

- For all protocols: microcentrifuge, disposable gloves, sterile pipet tips, sterile 1.5-2 ml tubes.
- For Agarose-Out protocol equipment necessary for visualization and excision of DNA band. In most cases: UV lamp and the scalpel, heating block or water bath set at 55°C.

# Protocol I

# PCR / DNA Clean-Up

The Protocol allows for quick purification of PCR products, restriction fragments, DNA molecules after enzymatic treatment and chemical or isotopic labeling.

- 1. Apply 30 μl of activation **Buffer Uni** onto the spin-column (do not spin) and keep it at room temperature till transfering mixture (p. 3) to the spin-column.
  - Addition of Buffer Uni onto the center of the resin enables complete wetting of membranes and maximal binding of DNA.
  - The membrane activation should be done before starting isolation procedure. The minimum activation time is 5-15 min.
- 2. Add 2 volumes of **Basic** buffer to 1 volume of the to DNA sample and mix.
  - For example, add 200 μl of Basic buffer to 100 μl DNA sample.
  - ο Maximum volume of a DNA sample can not exceed 200 μl. The minimum volume of DNA sample is 40 μl. If the sample volume is less than 40 μl, bring to a volume of 40 μl with sterile distilled water.
- 3. Apply the mixture to the **DNA binding spin-column** and centrifuge at 11 000 x g for 1 min. Remove the spin-column, pour off supernatant and place back into the receiver tube.
- 4. Add 500 μl of **Wash UX1** buffer and spin down at 11 000 x g for 1 min.
- 5. Remove spin column, pour the supernatant off, replace back the spin-column.
- 6. Add 600 μl of **Wash UX2** buffer and spin down at 11 000 x g for 1 min.
- 7. Remove spin column, pour the supernatant off, replace back the spin-column.
- 8. Spin down at 11 000 x g for 1 min to remove traces of **Wash UX2** buffer.

- Place spin-column into new receiver tube (1.5-2 ml). Add 50-150 μl of Elution buffer to elute bound DNA.
  - Addition of eluting buffer directly onto the center of the membrane improves DNA yield.
  - To improve recovery of larger DNA fragments (above 5 kb) it is recommended to elute with buffer heated to 80°C.
  - **o** For elution of DNA the Elution buffer is highly recommended. The buffer is prepared using ultrapure water with trace addition of buffering compounds. The Elution buffer will not interfere with subsequent DNA manipulations, such as DNA sequencing, ligation or restriction digestion, among others.
  - $\circ$  It is possible to reduce the volume of eluting buffer below 50  $\mu$ l (no less than 20  $\mu$ l). However, recovery of DNA will gradually decrease.
- 10. Incubate spin-column/receiver tube assembly for 1 min at room temperature.
- 11. Spin down at 11 000 x g for 1 min.
- Remove spin column, cap the receiver tube. Isolated DNA is ready for analysis/ manipulations. It can be stored at 2-8°C or (preferred) at -20°C.

# Protocol II

# Agarose-Out

Protocol is designed to isolate ultrapure linear or circular DNA molecules, ranging in size from approximately 100 bp to 10 kb, from TAE- or TBE-agarose gels.

- 1. Apply 30 μl of activation **Buffer Uni** onto the spin-column (do not spin) and keep it at room temperature till transfering dissolved agarose solution to the spin-column.
  - Addition of Buffer Uni onto the center of the resin enables complete wetting of membranes and maximal binding of DNA.
  - The membrane activation should be done before starting isolation procedure. The minimum activation time is 5-15 min.
- Cut out DNA from agarose gel. Weight the agarose piece, the weight should not exceed 250 mg. Place the agarose slice(s) into Eppendorf tube.
  - Avoid excess of agarose cut out along with DNA fragment.
  - If weight of the gel slice exceeds 250 mg, use more than one column.

- **o** It is highly recommended that electrophoresis buffer is not re-used, due to pH changes which negatively affect DNA isolation from a gel.
- 3. Add 2.5 volumes of **Basic** buffer to 1 volume of gel (100 mg ~ 100 μl). Mix by three-fold inverting.
  - For example, add 250 μl of Orange A buffer to each 100 mg of gel.
- Incubate in heating block or water bath at 55°C, mixing every 1–2 min by two-fold inverting, until agarose will dissolve completely.
  - Agarose slice(s) will dissolve within 5 to 10 min, depending on used agarose gel concentration and weight of agarose piece.
- 5. Apply up to  $600 \mu l$  of a dissolved agarose solution to the **DNA binding spin-column** and centrifuge at  $11\,000 \, x \, g$  for 1 min. Remove the spin-column, pour off supernatant and place back into the receiver tube.
- Transfer the remaining mixture to the same <u>DNA binding spin-column</u> and centrifuge at 11 000 x g for 1 min. Remove the spin-column, pour off supernatant and place back into the receiver tube.
- 7. Add 400 µl of **Basic** buffer and spin down at 11 000 x g for 1 min.
- 8. Remove spin column, pour off supernatant, replace back the spin-column.
- 9. Add 600 μl of **Wash UX2** buffer and spin down at 11 000 x g for 1 min.
- 10. Remove spin column, pour off supernatant, replace back the spin-column.
- 11. Spin down at 11 000 x g for 1 min to remove traces of Wash UX2 buffer.
- 12. Place spin-column into new receiver tube (1.5–2 ml) and add 50–80 μl of **Elution** buffer to elute bound DNA.
  - Addition of eluting buffer directly onto the center of the membrane improves DNA yield.
  - To improve recovery of larger DNA fragments (above 5 kb) it is recommended to elute with buffer heated to 80°C.
  - For elution of DNA the Elution buffer is highly recommended. The buffer is prepared using ultrapure water with trace addition of buffering compounds. The Elution buffer will not interfere with subsequent DNA manipulations, such as DNA sequencing, ligation or restriction digestion, among others.
  - $\circ$  It is possible to reduce the volume of eluting buffer below 50  $\mu$ l (no less than 20  $\mu$ l). However, recovery of DNA will gradually decrease.
  - To obtain optimal results of automated DNA sequencing it is recommended to use 0.3-0.6 pmols of DNA template per reaction (examplified by 200-400 ng of 1 kb DNA fragment).

- 13. Incubate spin-column/receiver tube assembly for 1 min at room temperature.
- **14**. Spin down at 11 000 x g for 1 min.
- 15. Remove spin column, cap the receiver tube. Isolated DNA is ready for analysis/manipulations. It can be stored at 2-8°C or (preferred) at -20°C.

# Protocol III

## **Plasmid Isolation**

The Protocol allows for quick isolation of plasmid DNA from bacterial cultures.

- Apply 30 μl of activation <u>Buffer Uni</u> onto the spin-column (do not spin) and keep it at room temperature till transfering lysate to the spin-column.
  - Addition of Buffer Uni onto the center of the resin enables complete wetting of membranes and maximal binding of DNA.
  - **o** The membrane activation should be done before starting isolation procedure. The minimum activation time is 5-15 min.
- 2. Pour cells from overnight culture (11–14 h) into 1.5–2 ml tubes. Spin down 1.0–3.0 ml in a microcentrifuge at 12 000 x g for 2 min. Pour off the supernatant and blot tubes upside-down on paper towel to remove any remaining media.
  - **o** E.coli strains recommended for plasmid isolation have endA- genotype, such as: DH5a, DH1, JM103-109, XL1-Blue, MM294 and C600. While endA+ strains, such as BL21, RR1, DH11S, JM101, HB101,TG1 and TB1 can also be used, but they yield lower quality DNA.
- 3. Add 250  $\mu$ l of **Cell R** buffer and completely resuspend the cell pellet.
- Add 250 μl of blue-coloured Lysis Blue buffer. Mix gently, but completely by several-fold inverting, until uniform blue colour of cell resuspension is obtained.
  - Alkaline Lysis Blue buffer contains SDS, which can precipitate at temperatures below 20°C. In this case warm the buffer up in 37°C water bath, until clarified.
  - Forceful mixing should be avoided, as it can cause irreversible denaturation of plasmid DNA molecules as well as contamination with genomic DNA fragments.
- Add 350 μl of neutralization and binding buffer Neutral B. Mix by several-fold inverting, until blue colour will disappear.

- 6. Spin down in a microcentrifuge at 12 000 x g for 7 min.
- 7. Apply up to  $600 \mu l$  of a clear supernatant to the **DNA binding spin-column** and centrifuge at  $11\,000 x$  g for 1 min. Remove the spin-column, pour off supernatant and place back into the receiver tube.
- 8. Transfer the remaining mixture to the same **DNA binding spin-column** and centrifuge at 11 000 x g for 1 min. Remove the spin-column, pour off supernatant and place back into the receiver tube.
- 9. Add 500 μl of **Wash UX1** buffer and spin down at 11 000 x g for 1 min.
- 10. Remove spin-column, pour off supernatant, replace back spin-column.
- 11. Add 600 μl of Wash UX2 buffer and spin down at 11 000 x g for 1 min.
- 12. Remove spin-column, pour off supernatant, replace spin-column.
- 13. Spin down at 11 000 x g for 1 min to remove traces of the Wash UX2 buffer.
- Place spin-column into new receiver tube (1.5-2 ml). Add 50-100 μl of Elution buffer to elute bound DNA.
  - Addition of eluting buffer directly onto the center of the membrane improves DNA yield.
  - f O To improve recovery of larger plasmids (above 6 kb) it is recommended to elute with buffer heated to 80°C.
  - **o** For elution of DNA the Elution buffer is highly recommended. The buffer is prepared using ultrapure water with trace addition of buffering compounds. The Elution buffer will not interfere with subsequent DNA manipulations, such as DNA sequencing, ligation or restriction digestion, among others.
  - $\circ$  It is possible to reduce the volume of eluting buffer below 50  $\mu$ l (no less than 20  $\mu$ l). However, recovery of DNA will gradually decrease.
- 15. Incubate spin-column/receiver tube assembly for 1 min at room temperature.
- **16.** Spin down at 11 000 x g for 1 min.
- Remove spin-column, cap the receiver tube. Plasmid DNA is ready for analysis/ manipulations. It can be stored either at 2-8°C or (preferred) at -20°C.

# **Safety Information**

### **Buffer Uni**

#### Danger



**H314** Causes severe skin burns and eye damage.

**P280** Wear protective gloves/protective clothing/eye protection/face protection.

**P301+P330+P331** If swallowed: Rinse mouth. Do not induce vomiting.

P303+P361+P353 If on skin (or hair): take off immediately all contaminated clothing. Rinse skin with water [or shower].

P305+P351+P338 If in eyes: rinse cautiously with water for several minutes. Remove contact lenses, if present and easy to do. Continue rinsing.

**P310** Immediately call a poison center/doctor. **P405** Store locked up.

### **Basic**

### Warning



H302+H332 Harmful if swallowed or if

**H412** Harmful to aquatic life with long lasting effects.

P273 Avoid release to the environment.
P301+P312 If swallowed: call a poison

center/ doctor/... if you feel unwell.

P304+P340 If inhaled: remove person to

P304+P340 If inhaled: remove person to fresh air and keep comfortable for breathing. EUH032 Contact with acids liberates very toxic gas.

### Lvsis Blue

#### Warning



H315 Causes skin irritation.

H319 Causes serious eye irritation.

**P280** Wear protective gloves/protective clothing/eve protection/face protection.

P305+P351+P338 If in eyes: rinse cautiously with water for several minutes. Remove contact lenses, if present and easy to do. Continue rinsing.

**P337+P313** If eye irritation persists: get medical advice/ attention.

P302+P352 If on skin: wash with plenty of water.

P332+P313 If skin irritation occurs: get medical advice/attention.

### **Neutral B**

#### )anger



H302+H332 Harmful if swallowed or if inhaled.

H315 Causes skin irritation. H319 Causes serious eye irritation.



**H334** May cause allergy or asthma symptoms or breathing difficulties if inhaled.

H317 May cause an allergic skin reaction.

P280 Wear protective gloves/protective

clothing/eye protection/face protection. **P284** [In case of inadequate ventilation] wear respiratory protection.

**P301+P312** If swallowed: call a poison center/ doctor if you feel unwell.

P304+P340 If inhaled: remove person to fresh air and keep comfortable for breathing. P305+P351+P338 If in eyes: rinse cautiously with water for several minutes. Remove contact lenses, if present and easy to do. Continue rinsing.

**P333+P313** If skin irritation or rash occurs: get medical advice/attention.

### Wash UX1

### Warning



H226 Flammable liquid and vapour. H302+H332 Harmful if swallowed or if inhaled.

H315 Causes skin irritation.



H319 Causes serious eye irritation.
P210 Keep away from heat, hot surfaces, sparks, open flames and other ignition sources. No smoking.

**P280** Wear protective gloves/protective clothing/eye protection/face protection.

**P301+P312** If swallowed: call a poison center/ doctor if you feel unwell.

P302+P352 If on skin: wash with plenty of water.

P304+P340 If inhaled: remove person to fresh air and keep comfortable for breathing. P305+P351+P338 If in eyes: rinse cautiously with water for several minutes. Remove contact lenses, if present and easy to do. Continue rinsing.



### Danger



 $\ensuremath{\textbf{H225}}$  Highly flammable liquid and vapour.

H319 Causes serious eye irritation.

**P210** Keep away from heat, hot surfaces, sparks, open flames and other ignition sources. No smoking.

**P280** Wear protective gloves/protective clothing/eye protection/face protection.

P305+P351+P338 If in eyes: rinse cautiously with water for several minutes. Remove contact lenses, if present and easy to do. Continue rinsing.

**P403+P235** Store in a well-ventilated place. Keep cool.

**P337+P313** If eye irritation persists: get medical advice/ attention.

# • GeneMATRIX is synthetic, new generation DNA- and RNA-binding membrane, selectively binding nucleic acids to composite silica structures.

Novel binding and washing buffers are developed to take full advantage of GeneMATRIX capacity, yielding biologically active, high-quality nucleic acids. Matrix is conveniently pre-packed in ready-to-use spin-format. Unique chemical composition of the matrixes along with optimized construction of spin-columns improve the quality of final DNA or RNA preparation. To speed up and simplify isolation procedure, the key buffers are colour coded, which allows monitoring of complete solution mixing and makes purification procedure more reproducible.

As a result, we offer kits, containing matrixes and buffers that guarantee rapid, convenient, safe and efficient isolation of ultrapure nucleic acids. Such DNA or RNA can be directly used in subsequent molecular biology applications, such as: restriction digestion, dephosphorylation, kinasing, ligation, protein-DNA interaction studies, sequencing, blotting, in vitro translation, cDNA synthesis, hybrydization among others. Additional advantage is reproducibility of matrix performance, as component preparation is carried at Eurx Ltd.

GeneMATRIX Basic DNA Purification Kit is a useful basic tool in any laboratory working with DNA. It allows to perform three basic laboratory techniques: DNA purification after enzymatic treatment, isolation of DNA from agarose gels and the isolation of plasmid DNA from bacterial cultures. Combining these capabilities kit allows to streamlin work and minimizing the costs of research projects.

PCR / DNA Clean-Up Protocol allows for purification of DNA fragments, which were subjected to or obtained as a result of various modifications and reactions: PCR products, restriction digests, after kinasing, dephosphorylation, end-trimming/repair, ligation, enzymatic or chemical modification, among others. Fragment of sizes from approximately 100 bp to over 15 kb can be obtained in ultrapure form. Effectively removed are contaminants such as: ethidum bromide, primers (below 40 nt), short double-stranded DNA (below 20 bp), RNA, Tag DNA Polymerase, Pfu DNA Polymerase, endo- and exonucleases, DNA-binding and modifying proteins, BSA and other enzymes/proteins, lipids, endotoxins, dyes, detergents, nucleotides, radio- and chemical labels, EDTA, problematic restriction and ligation inhibitors, buffers and salts.

Agarose-Out Protocol is designed to isolate linear or circular DNA molecules, ranging in size from approximately 100 bp to 10 kb, from TAE- or TBE-agarose gels. It is also possible to purify DNA fragments up to 20 kb or more, with gradually decreasing yields. Besides agarose many other contaminants are effectively removed: ethidium bromide, RNA, primers, enzymes and other proteins, lipids, endotoxins, dyes,

detergents, nucleotides, radio- and chemical labels, EDTA, problematic restriction and ligation inhibitors, buffers and salts.

Plasmid Isolation Protocol allows for purification of high purity plasmid DNA from various species of bacteria, including recombinant Escherichia coli strains. Plasmid DNA contaminants such as: RNA, single-stranded DNA, enzymes/proteins, lipids, dyes, detergents, nucleotides, EDTA, problematic restriction and ligation inhibitors, buffers and salts are effectively removed from crude bacterial lysate. Coloured lysis buffer helps both in monitoring cell solubilization progress as well as simultaneous processing of multiple samples.

For all protocols optimized buffer is added to provide selective conditions for DNA binding to the GeneMATRIX membranes during brief centrifugation, while contaminants pass through the spin-column. Traces of contaminants remaining on the membrane are efficiently removed in two wash steps. High-quality DNA is then eluted in low salt buffer, e.g.: Tris-HCl, TE or water. Isolated DNA is ready for downstream applications without the need for ethanol precipitation.



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