

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.

● **Cat. no. E3545**

EURx Ltd. 80-297 Gdansk Poland
ul. Przyrodników 3, NIP 957-07-05-191
KRS 0000202039, www.eurx.com.pl
orders: email: orders@eurx.com.pl
tel. +48 58 524 06 97, fax +48 58 341 74 23



Table of Contents

Introductory Notes.....	3
Equipment and reagents to be supplied by user:	4
Protocol I	4
PCR / DNA Clean-Up.....	4
Protocol II.....	5
Agarose-Out.....	5
Protocol III.....	7
Plasmid Isolation.....	7
Safety Information	9

Content	50 preps E3545-01	150 preps 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	

* 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 µg 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.

NOTE 5 • Elution buffer is a low salt solution, that contains no metal ion chealators (e.g. EDTA) that can inhibit subsequent enzymatic reactions. Elution buffer composition is suitable for downstream applications such as digestion with restriction enzymes, phosphorylation, ligation, Sanger sequencing, NGS etc. It is also possible to elute the DNA with Tris-HCl, water or TE.

Equipment and reagents to be supplied by user:

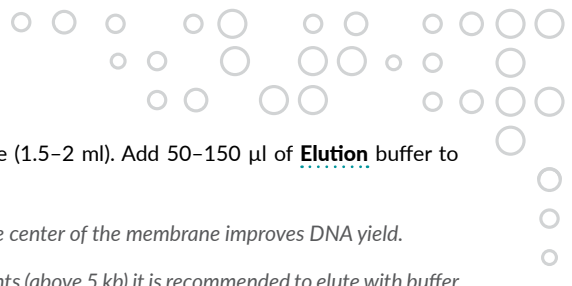
- For all protocols: microcentrifuge, disposable gloves, sterile pipet tips, sterile 1.5–2 ml tubes.
- For Agarose-Out protocol – equipment necessary for visualization and excision of DNA band. In most cases: UV lamp and the scalpel, heating block or water bath set at 55°C.

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 transferring 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 DNA sample and mix.
 - *For example, add 200 µl of Basic buffer to 100 µl DNA sample.*
 - *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, discard the flow-through and place back into the collection tube.
4. Add 500 µl of **Wash UX1** buffer and spin down at 11 000 x g for 1 min.
5. Remove the spin-column, discard the flow-through and place back into the collection tube.
6. Add 600 µl of **Wash UX2** buffer and spin down at 11 000 x g for 1 min.
7. Remove the spin-column, discard the flow-through and place back into the collection tube.
8. Spin down at 11 000 x g for 1 min to remove traces of **Wash UX2** buffer.


- 
9. 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.
 - 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.
 - It is possible to reduce the volume of eluting buffer below 50 μl (no less than 20 μl). 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 $\times g$ for 1 min.
 12. 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 transferring 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.
2. Excise the DNA fragment from the agarose gel. Place the agarose sample(s) into Eppendorf tube and weigh the gel slice. The weight per single preparation should not exceed 250 mg.
 - Minimize the size of the gel slice by removing excess agarose.
 - For the gel slices up to 250 mg, use 1.5 ml Eppendorf tube.

- 
- 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 Basic buffer to each 100 mg of gel.
 4. Incubate in heating block or water bath at 55°C, mixing every 1–2 min by two-fold inverting, until agarose has dissolved completely.
 - Agarose slice(s) will dissolve within 5 to 10 min, depending on agarose gel concentration and weight.
 5. Apply up to 600 µl 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, discard the flow-through and place back into the collection tube.
 6. 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, discard the flow-through and place back into the collection tube.
 7. Add 400 µl of **Basic** buffer and spin down at 11 000 x g for 1 min.
 8. Remove the spin-column, discard the flow-through and place back into the collection tube.
 9. Add 600 µl of **Wash UX2** buffer and spin down at 11 000 x g for 1 min.
 10. Remove the spin-column, discard the flow-through and place back into the collection tube.
 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.
 - It is possible to reduce the volume of eluting buffer below 50 µl (no less than 20 µl). 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 (exemplified 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.

1. Apply 30 µl of activation **Buffer Uni** onto the spin-column (do not spin) and keep it at room temperature till transferring 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.
 - The membrane activation should be done before starting isolation procedure. The minimum activation time is 5-15 min.
2. Pellet 1-3 ml of the overnight culture in 1.5-2 ml Eppendorf tubes by centrifuging 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.
 - *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 µl of **Cell R** buffer and resuspend the cell pellet.
 - The bacteria should be resuspended completely by vortexing or pipetting until no aggregates are visible.
4. Add 250 µl of blue-coloured **Lysis Blue** buffer. Mix gently by several-fold inverting, until uniform blue colour of cell lysate is visible.
 - The solution becomes viscous and slightly clear.
 - 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.

- 
5. Add 350 µl of neutralization and binding buffer **Neutral B**. Mix by several-fold inverting, until blue colour disappear.
 6. Spin down at 12 000 x g for 7 min.
 7. Apply up to 600 µl of a clear supernatant to the **DNA binding spin-column** and centrifuge at 11 000 x g for 1 min. Remove the spin-column, discard the flow-through and place back into the collection 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, discard the flow-through and place back into the collection tube.
 9. Add 500 µl of **Wash UX1** buffer and spin down at 11 000 x g for 1 min.
 10. Remove the spin-column, discard the flow-through and place back into the collection tube.
 11. Add 600 µl of **Wash UX2** buffer and spin down at 11 000 x g for 1 min.
 12. Remove the spin-column, discard the flow-through and place back into the collection tube.
 13. Spin down at 11 000 x g for 1 min to remove traces of the **Wash UX2** buffer.
 14. 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.
 - To improve recovery of larger plasmids (above 6 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.
 - It is possible to reduce the volume of eluting buffer below 50 µl (no less than 20 µl). 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.
 17. 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 inhaled.

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 fresh air and keep comfortable for breathing.

EUH032 Contact with acids liberates very toxic gas.

Lysis Blue

Warning



H315 Causes skin irritation.

H319 Causes serious eye irritation.

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.

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

Danger



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.

Wash UX2

Danger



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.

○ **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: ethidium bromide, primers (below 40 nt), short double-stranded DNA (below 20 bp), RNA, Taq 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,

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, hybridization among others. Additional advantage is reproducibility of matrix performance, as component preparation is carried at Eurx Ltd.

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.

EURx Ltd. 80-297 Gdansk Poland
ul. Przyrodnikow 3, NIP 957-07-05-191
KRS 0000202039, www.eurx.com.pl
orders: email: orders@eurx.com.pl
tel. +48 58 524 06 97, fax +48 58 341 74 23

