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Product Information

GenElute™ Direct mRNA Miniprep Kit

Catalog Numbers DMN10, DMN 70

TECHNICAL BULLETIN

Product Description

The GenElute[™] Direct mRNA Miniprep kit provides a convenient format to isolate polyadenylated mRNA directly from mammalian cells and tissues. The direct mRNA isolation procedure is based on that of Badley.¹ Up to 10^7 mammalian cells or 40 mg tissue are lysed and homogenized, either with the filtration columns provided or with a mechanical homogenizer. RNase is eliminated during a 10 minute proteinase K digestion. Sodium chloride is added, and polyadenylated RNA is captured on oligo(dT) polystyrene beads during a 10 minute incubation. For further enrichment, RNA may be released from the beads into fresh lysis solution and recaptured with the original beads. After 3 washes in a spin column, purified mRNA is eluted in 100 µl of 10 mM Tris-HCl, pH 7.4. Note that if all traces of DNA contamination must be eliminated for RT-PCR, further treatment with Amplification Grade DNase I is recommended, Catalog No. AMPD1.

Precautions and Disclaimer

The GenElute Direct mRNA Miniprep Kit is for laboratory use only. Not for drug, household or other uses. Wear gloves, safety glasses, and suitable protective clothing when handling any reagent provided with the kit. See the Material Safety Data Sheet (MSDS).

RNases are ubiquitous and very stable proteins, which are concerns for any researcher attempting to isolate RNA. Conditions during lysis inhibit RNase activity, and the proteinase K digestion eliminates RNases present in the original cell or tissue sample. **Care must be taken not to introduce RNase, especially during the wash and elution steps.** Use RNase-free pipette tips, preferably those having an aerosol barrier. Wear latex, nitrile or vinyl gloves and change them frequently. Keep bottles and tubes closed when not in use. References 2 - 4 at the end of this bulletin are good sources of information on working with RNA.

| Reagents Provided | Catalog No. | DMN10 10 Preps | DMN70 70 Preps |
|--|-------------|-------------------|-------------------|
| Lysis Solution | L2163 | 20 ml | 120 ml |
| Proteinase K | P2308 | 5 mg | 25 mg |
| 40% Glycerol | P0484 | 0.6 ml | 3 ml |
| 5 M NaCl Solution | S5150 | 1.5 ml | 8 ml |
| Oligo(dT) Polystyrene Beads | O8512 | 0.3 ml | 2 ml |
| Wash Solution | W2140 | 4 ml | 30 ml |
| Low Salt Wash Solution | W2265 | 8 ml | 60 ml |
| Elution Solution (10 mM Tris-HCl, pH 7.4) | E8026 | 1.5 ml | 10 ml |
| GenElute Filtration Columns | CP9346* | 10 each | 70 each |
| Spin Filters with Tubes | C9596 | 10 each | 70 each |
| Dolphin Collection Tubes | D7688 | 10 each | 70 each |
| Collection Tubes, 2.0 ml capacity | T5449 | 10 each | 70 each |

*MilliporeSigma continually seeks ways to improve our products. Please note that the product code for the GenElute Filtration Columns has changed from C9346 to CP9346. This change has been made to streamline and make more consistent, all the GenElute products across the line. The performance and functionality of the C9346 and CP9346 filtration columns are equivalent.

Equipment and Reagents Required But Not Provided

- Heating block set at 65 °C
- Tissue homogenizer and vessels for tissue samples
- Microcentrifuge capable of 16,000 x g (maximum speed in rotors that hold 18 tubes)

Storage

Store at room temperature. If any reagent forms a precipitate, see Preparation Instructions that follow.

Preparation Instructions

Before beginning the procedure, do the following:

- Thoroughly mix reagents. Examine for precipitation. If any reagent forms a precipitate, warm at 55-65 °C until the precipitate dissolves and allow to cool to room temperature before use.
- Dissolve the proteinase K (P2308) in the 40% glycerol (P0484). Use 0.5 ml of 40% glycerol for 5 mg proteinase K (10 prep package) or 2.5 ml of 40% glycerol for 25 mg proteinase K (70 prep package). Let sit at room temperature for a few minutes, and mix thoroughly before use. This product is stable at room temperature in its dry form, but store the proteinase K solution at 2-8 °C.
- 3. Add the proteinase K solution to a sufficient volume of Lysis Solution **(L2163)** for that day's use. The amount of Lysis Solution containing proteinase K required per preparation is 0.5 ml for cultured cells and 1.0 ml for tissue. Add 20 μ l of proteinase K solution per ml of Lysis Solution and mix thoroughly.
- Transfer approximately 120 μl of Elution Solution (E8026) per preparation into a microcentrifuge tube and heat to 65 °C in a heating block.
- 5. Ensure that the oligo(dT) beads **(O8512)** are at room temperature and vortex thoroughly before use.

Procedure

Note: All centrifugation speeds are given in units of g. Please refer to Table 1 for information on converting g-force to rpm. If centrifuges/rotors for the required g-forces are not available, use the maximum g-force possible and increase the spin time proportionally. Spin until all liquid passes through the column.

Table 1. Conversion of Centrifugal Force (in units of g) to RPM for Common Rotors

| Centrifuge | Rotor | Tubes (max) | Radius (cm) | RPM at 300 x <i>g</i> | RPM at 16,000 x <i>g</i> |
|-------------------|-----------|----------------|----------------|--------------------------|-----------------------------|
| Eppendorf 5410 | - | 12 | 5.8 | 2,143 | 15,652 |
| 5415C | F45-18-11 | 18 | 7.3 | 1,917 | 14,000 |
| 5415D&R | F45-24-11 | 24 | 8.3 | 1,801 | 13,155 |
| 5417C,D,&R | F45-30-11 | 30 | 9.5 | 1,681 | 12,279 |

See table above for spin speeds in rpm for selected common centrifuges and rotors. The correct rpm for unlisted rotors can be calculated using the formula:

$$RPM = \sqrt{RCF / 1.118 \times 10^{-5}} r$$

where RCF = required gravitational acceleration (relative centrifugal force) in units of *g*; *r* = radius of the rotor in cm; RPM = the number of revolutions per minute required to achieve the necessary *g*-force.

All steps are carried out at room temperature except as noted. All centrifugation (spin) steps should be performed at maximum speed (16,000 x g).

- 1. **Harvest cells or prepare tissue.** For best yields of intact RNA, use only rapidly growing cells before they reach their maximum density or harvest tissue immediately from a freshly sacrificed animal.
 - a. <u>Suspension cell cultures</u>: Count cells. Pellet up to 10⁷ cells for 5 minutes at 300 x g.
 Remove the culture medium completely and discard. Continue with step 2.
 - b. <u>Attached cell cultures</u>: Release attached cells with trypsin and pellet before lysis. To release cells with trypsin, see the product information sheet for **T3924** online for a typical use protocol. Count cells. Pellet up to 10⁷ cells and proceed as for suspension cultures.
 - c. <u>Mammalian tissue</u>: Quickly slice and weigh a piece of fresh or frozen tissue. Use 10 to 40 mg per preparation. Transfer to an appropriate vessel for homogenization and continue with step 2.

Note: Tissue may be flash-frozen in liquid nitrogen and stored at –70 °C for several months before preparing RNA. However, do not allow frozen tissue to thaw before disruption in Lysis Solution.

- Lyse and homogenize cells/tissue. This is a critical step that must be accomplished quickly and thoroughly.
 - a. <u>Pelleted cells</u>: Vortex the pellet to loosen the cells. Add 0.5 ml Lysis Solution containing proteinase K. Vortex or pipette thoroughly until all clumps disappear. Assemble a blue filtration column (CP9346) with a 2mL collection tube (T5449). Transfer the lysed cells into the filtration column. Centrifuge at maximum speed for 2 minutes. Discard the blue filtration column, and continue to step 3 with the homogenized lysate (flow-through liquid).
 - Mammalian tissue: Add 1 ml of Lysis Solution containing proteinase K and homogenize immediately until no visible pieces remain. Continue with step 3.

Note: Approximately 30 to 60 seconds of homogenization with a Brinkman Polytron PT 1200 is sufficient for most tissues. However, with fibrous tissues, such as muscle, several 30 second bursts are required to completely disrupt the tissue.

- 3. **Digest with proteinase K.** Incubate homogenized lysate at 65 °C for 10 minutes to degrade nucleases and other proteins. After incubation, remove the tube(s) from the heating block, and proceed with steps 4 and 5.
- 4. **Prepare mRNA for binding**. Add 32μ l of the 5 M NaCl solution **(S5150)** to the digested cell lysate, or 64 μ l of 5 M NaCl to the digested tissue lysate.

Note: The sodium chloride and oligo(dT) beads (step 5) may be added while the lysate is still warm.

- 5. Bind mRNA to oligo(dT) beads. Mix the oligo (dT) beads (O8512) thoroughly by vortexing and inverting until resuspended and homogenous. Add 25 μl of the resuspended oligo(dT) beads to the lysate-NaCl mixture, cap tube, and mix thoroughly by vortexing. Incubate the lysate/bead mixture at room temperature for 10 minutes. No mixing or rocking is necessary. During this incubation, the poly(A) tails of mRNA will hybridize with the oligo(dT) on the beads.
- Collect bead:mRNA complex. Pellet the oligo(dT) beads:mRNA complex by centrifuging for 5 minutes at maximum speed. Carefully remove and discard the supernatant, leaving behind approximately 50 μl to avoid disturbing the pellet.
- 7. Release and rebind mRNA (optional). For a more highly enriched mRNA preparation, bound material may be released from the beads into fresh lysis solution: mRNA is then rebound to the same beads. Add 0.5 ml of Lysis Solution and 32 µl of 5 M NaCl solution. Vortex thoroughly to resuspend the pellet. A series of short bursts on a vortex mixer are usually more effective than one long burst. Invert tube and inspect to verify complete release of pellet. Incubate suspension at 65 °C for 5 minutes. Remove from heat and incubate at room temperature for 5 minutes. Repellet the bead:mRNA complex for 2 minutes. Remove and discard all but ~50 μ l of the supernatant as above.

Note: This step may be omitted if rRNA contamination is not a concern, or if purification will be repeated as described in the note at the end of the procedure.

- 8. First wash. Resuspend the pellet in 350 μl of Wash Solution (W2140) by vortexing or pipetting. A series of short bursts on a vortex mixer will help release the pellet. Transfer the suspension into a GenElute spin filter-collection tube assembly (C9596) by pipetting. Make sure that all suspension is expelled from the pipette tip and that the pellet is completely removed from tube. Spin for 1-2 minutes at maximum speed. Remove the spin filter, discard the flow-through liquid, then place the spin filter back into the same collection tube.
- Second wash. Pipette 350 µl of Low Salt Wash Solution (W2265) into the column. Spin for 1-2 minutes. Spin for 1-2 minutes at maximum speed. Remove the spin filter, discard the flow-through liquid, then place the spin filter back into the same collection tube.
- Third wash. Pipette another 350 μl of Low Salt Wash Solution (W2265) into the column. Spin for 1-2 minutes at maximum speed.
- First elution. Transfer the spin filter into a fresh collection tube (T7688). Discard the flow-through liquid and the original collection tube. Pipette 50 μl of preheated (65 °C) Elution Solution (E8026) onto the spin filter ensuring that it contacts the bead:mRNA complex. Incubate for 2-5 minutes at 65 °C. Spin for 1-2 minutes at maximum speed. Save the flow-through liquid; it contains most of the purified mRNA.
- Second elution. To maximize recovery of mRNA, pipette an additional 50 μl of preheated Elution Solution onto the bead:mRNA complex. Incubate for 2-5 minutes at 65 °C. Spin for 1-2 minutes at maximum speed.

The poly(A)⁺ mRNA is now in the flow-through eluate (90-100 μ l total), and is ready for immediate use or storage at -70 °C. Keep the mRNA on ice whenever it is thawed for use.

Note: If a highly enriched preparation is desired, repurify the mRNA by adding 400 μ l of Lysis Solution, 32 μ l of 5 M NaCl solution, and a fresh 25 μ l aliquot of oligo(dT) beads to the 100 μ l eluted mRNA. Vortex and incubate 10 minutes at room temperature. Pellet the beads for 2 minutes, then wash and elute as in steps 8 through 12 above.

Results

Determining yield

Due to the low levels of mRNA present in cells and in tissues (only 1-5% of total RNA), estimating the concentration and quality of mRNA from minipreparations spectrophotometrically by measuring absorbance at 260 and 280 nm is not recommended. Typically, the absorbance must be read using the neat sample or with little dilution. As a consequence, most or all of the mRNA isolated will be used. The sample may be recovered for downstream applications if RNase-free cuvettes are used. However, fluorometric determination with the RiboGreen[®] RNA Quantitation Reagent from Molecular Probes is a more sensitive and practical method, and is recommended.

Expected yield

The yield of mRNA varies according to the organism and developmental stage. In general, younger and more rapidly growing cultures or animals will contain more RNA. Furthermore, RNA levels vary over a wide range in different tissues. Brain, heart, and skeletal muscle generally have low levels of RNA, while kidney, pancreas, spleen, and liver can contain up to 10-fold more RNA. Typical yields from 40 mg of rat liver or 10^7 HEK 293 cells are 5-10 µg of RNA by the RiboGreen assay method.

Ethanol precipitation

Some applications will require a more concentrated mRNA preparation than that prepared with this kit. The mRNA may be concentrated by adding 20 μ g of glycogen (Catalog No. G1767), 0.1 volume of 3 M sodium acetate buffer, pH 5.2 (Catalog No. S7899), and 3 volumes of ice cold absolute ethanol (Catalog No. E7023), and precipitating overnight at –20 °C. The mRNA can be stored indefinitely as an ethanol precipitate.

Before use, pellet the precipitated mRNA at 16,000 x g at 2-6 °C for 15 minutes. Carefully remove the supernatant, wash the pellet with 70% ethanol, and spin at the maximum speed for 3-5 minutes. Carefully remove the supernatant and allow the tube to air dry while lying flat with the lid open for 30 minutes. Once dry, the pellet can be resuspended in RNase-free molecular biology reagent water (Catalog No. W4502).

Agarose gel electrophoresis

The mRNA may be fractionated on a denaturing formaldehyde agarose gel as described by Farrell.³ Staining with SYBR[®] Green II (Catalog No. S9305) is recommended, because formaldehyde gels stained with SYBR Green II do not require destaining. The mRNA should appear as a series of very closely spaced bands or smear at 0.5 kb or greater. More intense bands at approximately 5.3 and 2.0 kb are residual 18S and 28S rRNA.

Northern analysis

The integrity of the mRNA can be determined by transferring the gel pattern to a nylon membrane and hybridizing with a probe specific for an mRNA present in the population. See Related Products list for products and procedures to perform Northern blots, label probes and perform hybridization. Intact mRNA should result in a discrete band or bands, with no smearing downwards.

RT-PCR

While most DNA is eliminated during mRNA isolation, no single procedure removes 100% of the DNA. Because PCR can detect even a single molecule of DNA, RNA samples should be digested with Amplification Grade DNase I (Catalog No. AMPD1) before RT-PCR, and parallel samples should be assayed without adding reverse transcriptase. These precautions are especially recommended if PCR primers do not span an intron, or if pseudogenes that lack the intron may be present in the target cells or tissue.

References

- Badley, J.E., *et al.*, A simple, rapid method for the purification of poly(A)⁺ RNA. *BioTechniques* 6(2), 114-116 (1988)
- Ausubel, F.M., et al., Current Protocols in Molecular Biology (John Wiley & Sons, NY, 1995) sections 4.1-4.10
- Farrell, Robert E., Jr. *RNA Methodologies*, 2nd Edition (Academic Press, NY, 1998) pp. 37-53 and 153-156 (Product Code Z35,035-4)
- Sambrook, J., et al., Molecular Cloning: A Laboratory Manual, 2nd ed., (Cold Spring Harbor Laboratory Press, Plainview, NY, 1989) pp. 7.3-7.8

Troubleshooting Guide

| Problem | Cause | Solution |
|---------------------|--|---|
| Clogged spin filter | Sample size was too large | For future preparations, use fewer cells or smaller tissue samples. Alternatively, undigested material may be pelleted for 2-5 minutes after the proteinase K digestion in step 3, and the supernatant transferred to a new tube before adding the 5 M NaCl solution and oligo(dT) beads. To salvage the current preparation, spin longer than 1-2 minutes until solutions pass through the spin filter. The yield and purity of mRNA will likely be reduced. |
| | Homogenization was incomplete | Cell lysates must be spun through filtration columns to shear the DNA. Tissues must be thoroughly homogenized until no visible particles remain. |
| | Digestion was incomplete | Store proteinase K at 2-8 °C after it is dissolved in glycerol. Add proteinase K solution to the Lysis Solution immediately before use. The enzyme is not stable in the Lysis Solution for extended storage. Verify that the homogenized cells or tissues were incubated at 65 °C for 10 minutes before the 5 M NaCl solution was added. |
| | Centrifugal force was low | If the microcentrifuge used cannot attain 16,000 x g, longer spin times may be required. |
| Low yield of mRNA | Cells or tissue had low mRNA levels | Yields will vary greatly among different types of cells and tissues. See "Expected Yield" in the Results section. |
| | Elution Solution was not pre-heated or samples were not incubated at 65 °C | Transfer ~120 μ l of Elution Solution per preparation into a microcentrifuge tube and heat to 65 °C in a heating block before starting the procedure. Incubate the bead:mRNA complex with Elution Solution for 2-5 minutes at 65 °C before spinning. |
| Degraded mRNA | Tissue or culture was too old | Use cultures before they reach maximum density or become fully confluent, and harvest tissues as rapidly as possible from freshly sacrificed animals. |
| | Cells or tissues were stored improperly | If immediate preparation of mRNA is not possible, flash-freeze cell pellets or small pieces of tissue in liquid nitrogen and store at –70 °C. Do not allow material to thaw before it is disrupted in Lysis Solution. |
| | Cells or tissue contained high levels of RNase | Cells such as monocytes and macrophages, and tissues such as pancreas, spleen, and thymus, are rich in RNases and require immediate and thorough disruption in Lysis Solution to prevent degradation of RNA. |
| | Cells or tissues were not disrupted sufficiently | Vortex or pipet cell lysates until no clumps remain. Homogenize tissues in Lysis Solution until no visible particles remain. |
| | Proteinase K digestion was incomplete | See "Digestion is incomplete" above. |
| | RNase was introduced during the procedure | Pay special attention to precautions for handling RNA samples and related lab equipment listed at the beginning of this bulletin and in references 2-4. |

Troubleshooting Guide (cont.)

| Problem | Cause | Solution |
|-------------------------------|---|--|
| Excessive rRNA contamination | Abundance of rRNA is high; sequence of rRNA contains poly(A) regions | Detectable amounts of rRNA are expected. Non-specific binding to oligo(dT) can occur due to the vast excess of rRNA over mRNA. Also, poly(A) regions in rRNA can bind specifically to the beads. If a more enriched preparation is desired, repurify the mRNA as described in the note under Step 12 of the Procedure. |
| | Miniprep capacity was exceeded | Repurify as described in the note under Step 12 of the Procedure. For future preparations, use smaller amounts of starting cells or tissue. |
| | Releasing and re- binding procedures were omitted | Repurify as described in the note under Step 12 of the Procedure. |
| Poor results in downstream | Salt was carried over into eluate | Spin beads dry before adding Elution Solution. |
| procedures | Improper storage or handling of mRNA | Store eluted mRNA in Elution Solution at -70 °C or as an ethanol precipitate at -70 °C until needed. Keep the mRNA on ice whenever it is thawed for use. |

| Related Products | Catalog No. | Related Products | Catalog No. |
|---|--------------------|--|----------------|
| GenElute™ mRNA Miniprep Kits | MRN10 and MRN70 | Deoxyribonuclease I, Amplification Grade | AMPD1 |
| Ethanol, Absolute, Molecular Biology Reagent | E7023 | Enhanced Avian HS RT-PCR Kit | HSRT100 |
| Glycogen, Molecular Biology Grade | G1767 | Enhanced Avian Reverse Transcriptase | A4464 |
| 3 M Sodium Acetate Buffer, pH 5.2, Molecular Biology Grade | S7899 | Deoxynucleotide (dNTP) Mix | D7295 |
| Agarose, Molecular Biology Grade | A9539 | JumpStart™ REDTaq [®] Readymix™ | P0982 |
| Formaldehyde, Molecular Biology Grade | F8775 | Taq DNA Polymerase | D1806 D4545 |
| MOPS-EDTA-Sodium Acetate Buffer | M5755 | PerfectHyb [™] Plus Hybridization Buffer | H7033 |
| RNA Sample Loading Buffer | R1386 | SYBR Green II, 10,000X concentrate | S9305 |
| RNA markers, 0.2-10 kb | R7020 | | |

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