
November 2020

RNeasy[®] Plus Micro Handbook

For purification of total RNA using gDNA Eliminator columns from small samples, including animal and human cells ($\leq 5 \times 10^5$), animal and human tissues (≤ 5 mg), microdissected cryosections and for RNA cleanup and concentration

Contents

Kit Contents	4
Storage	4
Intended Use	5
Safety Information.....	5
Quality Control.....	6
Introduction.....	7
Principle and procedure	8
Equipment and Reagents to Be Supplied by User	10
Important Notes.....	11
Determining the amount of starting material.....	11
Handling and storing starting material	13
Disrupting and homogenizing starting material	15
Carrier RNA	17
Limitations of small samples	18
Automated purification of RNA on QIAcube Instruments.....	19
Protocol: Purification of Total RNA from Animal and Human Cells	21
Protocol: Purification of Total RNA from Animal and Human Tissues	30
Protocol: Purification of Total RNA from Microdissected Cryosections.....	39
Troubleshooting Guide	45
Appendix A: General Remarks on Handling RNA.....	49
Appendix B: Storage, Quantification and Determination of Quality of RNA.....	52
Appendix C: RNA Cleanup and Concentration	56

Appendix D: Purification of Total RNA Containing Small RNAs from Cells.....	57
Appendix E: Acetone Precipitation of Protein from Lysates	60
Ordering Information	62
Document Revision History.....	66

Kit Contents

RNeasy Plus Micro Kit	(50)
Catalog no.	74034
Number of preps	50
gDNA Eliminator Mini Spin Columns (uncolored) (each in a 2 ml Collection Tube)	50
RNeasy MinElute® Spin Columns (pink) (each in a 2 ml Collection Tube)	50
Collection Tubes (1.5 ml)	50
Collection Tubes (2 ml)	100
Buffer RLT Plus*	45 ml
Buffer RW1*	45 ml
Buffer RPE† (concentrate)	11 ml
RNase-Free Water	3 x 10 ml
Carrier RNA, poly-A	310 µg
Quick-Start Protocol	1

* Contains a guanidine salt. Not compatible with disinfectants containing bleach. See page 5 for safety information.

† Before using for the first time, add 4 volumes of ethanol (96–100%) as indicated on the bottle to obtain a working solution.

Storage

The RNeasy Plus Micro Kit is shipped at ambient temperature. Store the RNeasy MinElute spin columns immediately upon receipt at 2–8°C. Store the remaining components of the kit dry at room temperature (15–25°C). All kit components are stable for at least 9 months under these conditions, if not otherwise stated on the label.

Intended Use


The RNeasy Plus Micro Kit is intended for research use. No claim or representation is intended to provide information for the diagnosis, prevention or treatment of a disease.

QIAcube® Connect is designed to perform fully automated purification of nucleic acids and proteins in molecular biology applications. The system is intended for use by professional users trained in molecular biological techniques and the operation of QIAcube Connect.

All due care and attention should be exercised in the handling of the products. We recommend all users of QIAGEN® products to adhere to the NIH guidelines that have been developed for recombinant DNA experiments or to other applicable guidelines.

Safety Information

When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, please consult the appropriate safety data sheets (SDSs). These are available online in convenient and compact PDF format at www.qiagen.com/safety where you can find, view and print the SDS for each QIAGEN kit and kit component.

<p>CAUTION</p> 	<p>CAUTION: DO NOT add bleach or acidic solutions directly to the sample-preparation waste.</p>
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Buffer RLT Plus contains guanidine thiocyanate and Buffer RW1 contains a small amount of guanidine thiocyanate. Guanidine salts can form highly reactive compounds when combined with bleach. If liquid containing these buffers is spilt, clean with suitable laboratory detergent

and water. If the spilt liquid contains potentially infectious agents, clean the affected area first with laboratory detergent and water and then with 1% (v/v) sodium hypochlorite.

Quality Control

In accordance with QIAGEN's ISO-certified Quality Management System, each lot of RNeasy Plus Micro Kit is tested against predetermined specifications to ensure consistent product quality.

Introduction

The RNeasy Plus Micro Kit is designed to purify total RNA from small amounts of cells (as little as 1 cell) or easy-to-lyse tissues. Suitable starting materials include laser-microdissected (LMD) cryosections, fine-needle aspirates (FNA) and FACS® sorted cells. Genomic DNA contamination is effectively removed using a specially designed gDNA Eliminator spin column. The purified RNA is ready to use and is ideally suited for downstream applications that are sensitive to low amounts of DNA contamination, such as quantitative, real-time RT-PCR.* The purified RNA can also be used in other applications, including:

- RT-PCR
- Differential display
- cDNA synthesis
- Northern, dot and slot blot analyses
- Primer extension
- Poly A+ RNA selection
- RNase/S1 nuclease protection
- Microarrays

The RNeasy Plus Micro Kit allows the parallel processing of multiple samples in less than 30 minutes. Methods involving the use of toxic substances, such as phenol and/or chloroform, or time-consuming and tedious methods, such as alcohol precipitation, are replaced by the RNeasy Plus procedure.

* Visit www.qiagen.com/geneXpression for information on standardized solutions for gene expression analysis from QIAGEN.

Principle and procedure

The RNeasy Plus Micro procedure integrates QIAGEN's technology for selective removal of double-stranded DNA with well-established RNeasy MinElute technology. Efficient purification of high-quality RNA in a small elution volume is guaranteed, without the need for additional DNase digestion.

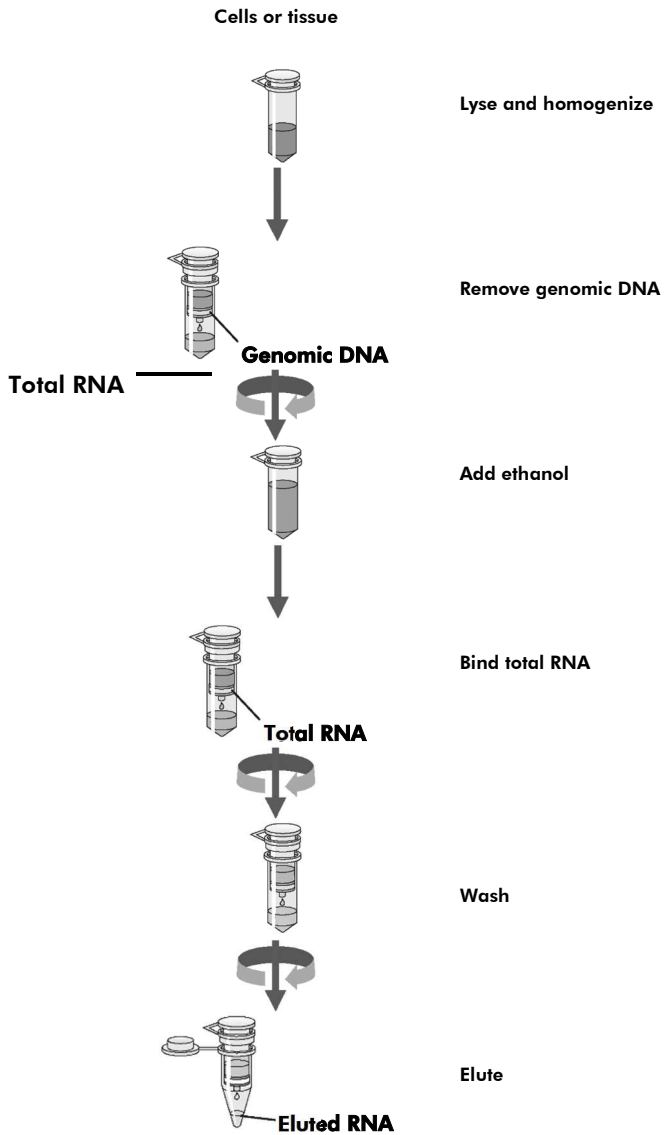
Biological samples are first lysed and homogenized in a highly denaturing guanidine-isothiocyanate-containing buffer, which immediately inactivates RNases to ensure isolation of intact RNA. The lysate is then passed through a gDNA Eliminator spin column. This column, in combination with the optimized high-salt buffer, efficiently removes genomic DNA.

Ethanol is added to the flow-through to provide appropriate binding conditions for RNA, and the sample is then applied to an RNeasy MinElute spin column, where total RNA binds to the membrane and contaminants are efficiently washed away. High-quality RNA is then eluted in as little as 14 µl of water.

With the RNeasy Plus Micro procedure, all RNA molecules longer than 200 nucleotides are purified. The procedure enriches for mRNA, since most RNAs <200 nucleotides (such as 5.8S rRNA, 5S rRNA and tRNAs, which together make up 15–20% of total RNA) are selectively excluded. The size distribution of the purified RNA is comparable to that obtained by centrifugation through a CsCl cushion, where small RNAs do not sediment efficiently. A modification of the RNeasy Plus Micro procedure for cells allows the purification of total RNA containing small RNAs, such as miRNA (see Appendix D, page 57).

In this handbook, different protocols are provided for different starting materials. The protocols differ primarily in the lysis and homogenization of the sample. Once the sample is applied to the gDNA Eliminator spin column, the protocols are similar (see flowchart, next page).

RNeasy Plus Micro Procedure



Equipment and Reagents to Be Supplied by User

When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, consult the appropriate safety data sheets (SDSs), available from the product supplier.

- 14.3 M β -mercaptoethanol (β -ME) (commercially available solutions are usually 14.3 M) or, alternatively, 2 M dithiothreitol (DTT) in water
- Ethanol (70% and 96–100%)*
- Sterile, RNase-free pipet tips
- Microcentrifuge (with rotor for 2 ml tubes)
- Vortexer
- Disposable gloves
- Reagent for RNA stabilization (see page 13):
 - For cell samples: RNAprotect[®] Cell Reagent[†] or liquid nitrogen
 - For tissue samples: RNAprotect Tissue Reagent[†] (stabilizes RNA only), Allprotect[®] Tissue Reagent[†] (stabilizes DNA, RNA and protein) or liquid nitrogen
- Equipment for sample disruption and homogenization (see pages 15–17). Depending on the method chosen, one or more of the following are required:
 - Trypsin and PBS
 - QIAshredder homogenizer[†]
 - Blunt-ended needle and syringe
 - Mortar and pestle
 - TissueRuptor II with TissueRuptor Disposable Probes[†]
 - TissueLyser II[†]

* Do not use denatured alcohol, which contains other substances, such as methanol or methylethylketone.

[†] For ordering information, see page 50.

Important Notes

Determining the amount of starting material

It is essential to use the correct amount of starting material to obtain optimal RNA yield and purity. The maximum amount that can be used is determined by:

- The type of sample and its RNA content
- The volume of Buffer RLT Plus required for efficient lysis
- The DNA removal capacity of the gDNA Eliminator spin column
- The RNA binding capacity of the RNeasy MinElute spin column

When processing samples containing high amounts of DNA or RNA, less than the maximum amount of starting material shown in Table 1 should be used, so that the DNA removal capacity of the gDNA Eliminator spin column and the RNA binding capacity of the RNeasy MinElute spin column are not exceeded.

When processing samples containing average or low amounts of RNA, the maximum amount of starting material shown in Table 1 can be used. However, even though the RNA binding capacity of the RNeasy MinElute spin column is not reached, the maximum amount of starting material must not be exceeded. Otherwise, lysis will be incomplete and cellular debris may interfere with the binding of RNA to the RNeasy spin column membrane, resulting in lower RNA yield and purity.

More information on using the correct amount of starting material is given in each protocol. Table 2 shows typical RNA yields from various cells and tissues.

Note: If the DNA removal capacity of the gDNA Eliminator spin column is exceeded, the purified RNA will be contaminated with DNA. Although the gDNA Eliminator spin column can bind more than 100 µg DNA, we recommend using samples containing less than 20 µg DNA

to ensure removal of virtually all genomic DNA. If the binding capacity of the RNeasy MinElute spin column is exceeded, RNA yields will not be consistent and may be reduced. If lysis of the starting material is incomplete, RNA yields will be lower than expected, even if the binding capacity of the RNeasy MinElute spin column is not exceeded.

Table 1. RNeasy MinElute spin column specifications

Maximum binding capacity	45 µg RNA
Maximum loading volume	700 µl
RNA size distribution	RNA >200 nucleotides*
Minimum elution volume	14 µl
Maximum amount of starting material	
Animal and human cells	5 x 10 ⁵
Animal and human tissues	5 mg

* Purification of total RNA containing small RNAs from cells is possible through a modification of the RNeasy Plus Micro procedure. For details, see Appendix D, page 57.

Table 2. Typical yields of total RNA with the RNeasy Plus Micro Kit

Sample type	Yield of total RNA* (µg)
Cell cultures (5 x 10⁵ cells)	
NIH/3T3	5
HeLa	7.5
COS-7	17.5
LMH	6
Huh	7.5
Mouse/rat tissues (5 mg)	
Embryo (13 day)	10
Brain	2–5
Kidney	10–15
Liver	20–30
Spleen	15–20
Thymus	20–25
Lung	5–10

* Amounts can vary due to factors, such as species, developmental stage and growth conditions. Since the RNeasy Plus Micro procedure enriches for mRNA and other RNA species >200 nucleotides, the total RNA yield does not include 5S rRNA, tRNA and other low-molecular-weight RNAs, which make up 15–20% of total cellular RNA.

Handling and storing starting material

Cells

After harvesting, cells should be immediately lysed in Buffer RLT Plus to prevent unwanted changes in the gene expression profile. This highly denaturing lysis buffer inactivates RNases and other proteins to prevent RNA degradation as well as downregulation or upregulation of transcripts.

If the cells are to be shipped to another lab for RNA purification, they should be pelleted, frozen in liquid nitrogen and transported on dry ice. Alternatively, the cells can be mixed with

RNAprotect Cell Reagent at room temperature (15–25°C) and then shipped at ambient temperature.

Tissues

RNA in harvested tissue is not protected until the sample is treated with RNAprotect Tissue Reagent, flash-frozen or disrupted and homogenized in the presence of RNase-inhibiting or denaturing reagents. Otherwise, unwanted changes in the gene expression profile will occur. It is therefore important that tissue samples are immediately frozen in liquid nitrogen and stored at –90°C to –65°C or immediately immersed in RNAprotect Tissue Reagent. An alternative to RNAprotect Tissue Reagent is Allprotect Tissue Reagent, which provides immediate stabilization of DNA, RNA and protein in tissue samples at room temperature (15–25°C).

The procedures for tissue harvesting and RNA protection should be carried out as quickly as possible. Frozen tissue samples should not be allowed to thaw during handling or weighing. After disruption and homogenization in Buffer RLT Plus (lysis buffer), samples can be stored at –90°C to –65°C for months.

Disrupting and homogenizing starting material

Efficient disruption and homogenization of the starting material is an absolute requirement for all total RNA purification procedures. Disruption and homogenization are 2 distinct steps.

- **Disruption:** Complete disruption of plasma membranes of cells and organelles is absolutely required to release all the RNA contained in the sample. Different samples require different methods to achieve complete disruption. Incomplete disruption results in significantly reduced RNA yields.
- **Homogenization:** Homogenization is necessary to reduce the viscosity of the lysates produced by disruption. Homogenization shears the high-molecular-weight genomic DNA and other high-molecular-weight cellular components to create a homogeneous lysate. Incomplete homogenization results in inefficient binding of RNA to the RNeasy MinElute spin column membrane and therefore significantly reduced RNA yields.

Some disruption methods simultaneously homogenize the sample, while others require an additional homogenization step. Table 3 gives an overview of different disruption and homogenization methods, and is followed by a detailed description of each method.

Table 3. Disruption and homogenization methods

Sample	Disruption method	Homogenization method
Microdissected samples	Addition of lysis buffer	Vortexing
Cells and fine-needle aspirates (FNA)	Addition of lysis buffer	TissueRuptor II or QIAshredder homogenizer or syringe and needle*
Tissues	TissueRuptor II	TissueRuptor II†
	Tissuelyser II‡	Tissuelyser II‡
	Mortar and pestle§	QIAshredder homogenizer or syringe and needle

* If processing $\leq 1 \times 10^5$ cells, the lysate can be homogenized by vortexing.

† Simultaneously disrupts and homogenizes individual samples.

‡ Simultaneously disrupts and homogenizes up to 192 samples in parallel. Results are comparable to those obtained using the TissueRuptor II or other rotor-stator homogenizer.

§ The TissueRuptor and Tissuelyser II usually give higher RNA yields than mortar and pestle.

Disruption and homogenization using the TissueRuptor II

The TissueRuptor II is a rotor–stator homogenizer that thoroughly disrupts and simultaneously homogenizes single tissue samples in the presence of lysis buffer in 15–90 seconds, depending on the toughness and size of the sample. The TissueRuptor II can also be used to homogenize cell lysates. The blade of the TissueRuptor disposable probe rotates at a very high speed, causing the sample to be disrupted and homogenized by a combination of turbulence and mechanical shearing. For guidelines on using the TissueRuptor II, refer to the *TissueRuptor II Handbook*. For other rotor–stator homogenizers, refer to suppliers' guidelines.

Disruption and homogenization using the Tissuelyser II

In bead-milling, tissues can be disrupted by rapid agitation in the presence of beads and lysis buffer. Disruption and simultaneous homogenization occur by the shearing and crushing action of the beads as they collide with the cells. The Tissuelyser II disrupts and homogenizes up to 48 tissue samples simultaneously when used in combination with the Tissuelyser Adapter Set 2 x 24, which holds 48 x 2 ml microcentrifuge tubes containing stainless steel beads of 5 mm mean diameter. The Tissuelyser II can also disrupt and homogenize up to 192 tissue samples simultaneously when used in combination with the Tissuelyser Adapter Set 2 x 96, which holds 192 x 1.2 ml microtubes containing stainless steel beads of 5 mm mean diameter. For guidelines on using the Tissuelyser II, refer to the *Tissuelyser II Handbook*. For other bead mills, refer to suppliers' guidelines.

Note: Tungsten carbide beads react with Buffer RLT Plus and must not be used to disrupt and homogenize tissues.

Disruption using a mortar and pestle

For disruption using a mortar and pestle, freeze the tissue sample immediately in liquid nitrogen and grind to a fine powder under liquid nitrogen. Transfer the suspension (tissue powder and liquid nitrogen) into a liquid-nitrogen-cooled, appropriately sized tube and allow the liquid nitrogen to evaporate without allowing the sample to thaw. Add lysis buffer and

continue as quickly as possible with the homogenization according to one of the 2 methods below.

Note: Grinding the sample using a mortar and pestle will disrupt the sample, but will not homogenize it. Homogenization must be performed afterwards.

Homogenization using QIAshredder homogenizers

Using QIAshredder homogenizers is a fast and efficient way to homogenize cell and tissue lysates without cross-contamination of samples. Up to 700 μ l of lysate is loaded onto a QIAshredder spin column placed in a 2 ml collection tube, and spun for 2 minutes at maximum speed in a microcentrifuge. The lysate is homogenized as it passes through the spin column.

Homogenization using a syringe and needle

Cell and tissue lysates can be homogenized using a syringe and needle. Lysate is passed through a 20-gauge (0.9 mm) needle attached to a sterile plastic syringe at least 5–10 times or until a homogeneous lysate is achieved. Increasing the volume of lysis buffer may be required to facilitate handling and minimize loss.

Carrier RNA

The RNeasy Plus Micro Kit contains poly-A RNA for use as carrier RNA. When added to lysates from very small samples, the carrier RNA may in some cases improve the recovery of total RNA. Carrier RNA is not required when processing more than 500 cells or more than about 2 μ g tissue.

As demonstrated in many different RT-PCR systems, the small amounts of poly-A RNA used as carrier RNA in total RNA purification do not interfere with subsequent RT-PCR, even when oligo-dT is used as a primer for reverse transcription. Reverse-transcription reactions typically contain an excess of oligo-dT primers, and the small amounts of poly-A used as carrier RNA are insignificant in comparison.

However, total RNA purified using poly-A RNA as carrier RNA is not compatible with protocols to amplify mRNA transcripts using oligo-dT primers. These include the Eberwine method and the QuantiTect® Whole Transcriptome Kit (the kit uses a mix of random and oligo-dT primers). For the Eberwine method, other types of RNA can be purchased separately for use as carrier RNA. Note, however, that tRNA and other RNAs <200 nucleotides will not bind to the RNeasy MinElute membrane and cannot be used as carrier RNA. For most applications, bacterial ribosomal RNA (e.g., from Roche, cat. no. 206938)* gives good results and can be used as an alternative to the poly-A RNA supplied with this kit. However, if amplifying mRNA transcripts with the QuantiTect Whole Transcriptome Kit, no carrier RNA of any type should be used in RNA purification.

Limitations of small samples

When purifying nucleic acids from particularly small samples (e.g., laser-microdissected samples), the amounts of RNA may be too small for quantification by spectrophotometry or even fluorometric assays. In this case, quantitative, real-time RT-PCR should be used for quantification.

When purifying RNA from less than 100 cells, stochastic problems with respect to copy number can occur. This is because some RNA transcripts may be present at very low copy numbers per cell, or only in a fraction of all cells in the sample of interest. For example, if a particular RNA transcript is present at an abundance of 1 copy per cell, and 10 cells are processed with RNA eluted in the recommended volume of 14 µl, there will be less than 1 copy of the transcript per microliter.

Whole transcriptome amplification can be carried out to generate sufficient amounts of RNA if several downstream assays need to be performed from a single small sample. However, care should be taken to include a sufficient amount of starting material in the amplification

* This is not a complete list of suppliers and does not include many important vendors of biological supplies. When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, consult the appropriate safety data sheets (SDSs), available from the product supplier.

reaction to avoid stochastic problems. The QuantiTect Whole Transcriptome Kit provides highly uniform amplification of the transcriptome.

Automated purification of RNA on QIAcube Instruments

Purification of RNA can be fully automated on QIAcube Connect or the classic QIAcube. The innovative QIAcube instruments use advanced technology to process QIAGEN spin columns, enabling seamless integration of automated, low-throughput sample prep into your laboratory workflow. Sample preparation using QIAcube instruments follows the same steps as the manual procedure (i.e., lyse, bind, wash and elute), enabling you to continue using the RNeasy Plus Micro Kit for purification of high-quality RNA.

QIAcube instruments are preinstalled with protocols for purification of plasmid DNA, genomic DNA, RNA, viral nucleic acids and proteins, plus DNA and RNA cleanup. The range of protocols available is continually expanding, and additional QIAGEN protocols can be downloaded free of charge at www.qiagen.com/qiacubeprotocols.



QIAcube Connect.

Protocol: Purification of Total RNA from Animal and Human Cells

Determining the correct amount of starting material

It is essential to use the correct amount of starting material to obtain optimal RNA yield and purity. The maximum amount depends on:

- The RNA content of the cell type
- The DNA removal capacity of the gDNA Eliminator spin column
- The RNA binding capacity of the RNeasy MinElute spin column (45 µg RNA)
- The volume of Buffer RLT Plus required for efficient lysis

In addition, cellular debris can reduce the binding capacity of the gDNA Eliminator and RNeasy MinElute spin columns. If processing a cell type not listed in Table 2 (page 13) and if there is no information about its RNA content, we recommend starting with no more than 5×10^5 cells.

Do not overload the gDNA Eliminator spin column, as this will lead to copurification of DNA with RNA. Do not overload the RNeasy MinElute spin column, as this will significantly reduce RNA yield and purity.

Counting cells is the most accurate way to quantitate the amount of starting material. As a guide, the number of HeLa cells obtained in various culture vessels after confluent growth is given in Table 4.

Important points before starting

- If using the RNeasy Plus Micro Kit for the first time, read “Important Notes” (page 11).
- If preparing RNA for the first time, read Appendix A (page 49).

- If using the TissueRuptor II, ensure that you are familiar with operating it by referring to the *TissueRuptor II User Manual* and *TissueRuptor II Handbook*.
- Cell pellets can be stored at -90°C to -65°C for later use or used directly in the procedure. Determine the number of cells before freezing. Frozen cell pellets should be thawed slightly so that they can be dislodged by flicking the tube in step 2. Homogenized cell lysates from step 3 can be stored at -90°C to -65°C for several months. Frozen lysates should be incubated at 37°C in a water bath until completely thawed and salts are dissolved. Avoid prolonged incubation, which may compromise RNA integrity. If any insoluble material is visible, centrifuge for 5 min at $3000\text{--}5000 \times g$. Transfer the supernatant to a new RNase-free glass or polypropylene tube, and continue with step 4.

Table 4. Growth area and number of HeLa cells in various culture vessels

Cell-culture vessel	Growth area (cm ²)*	Number of cells†
Multiwell plates		
96-well	0.32–0.6	$4\text{--}5 \times 10^4$
48-well	1	1×10^5
24-well	2	2.5×10^5
12-well	4	5×10^5
6-well	9.5	$1 \times 10^{6\dagger}$
Dishes		
35 mm	8	$1 \times 10^{6\dagger}$
Flasks		
40–50 ml	25	$3 \times 10^{6\dagger}$

* Per well, if multiwell plates are used; varies slightly depending on the supplier.

† Cell numbers are given for HeLa cells (approximate length = $15 \mu\text{m}$), assuming confluent growth. Numbers will vary for different kinds of animal and human cells, which vary in length from 10 to $30 \mu\text{m}$.

‡ This number of cells exceeds the maximum binding capacity of the RNeasy MinElute spin columns. To process this many cells, split the lysate into appropriate aliquots ($\leq 5 \times 10^5$ cells each) and load them onto separate RNeasy MinElute spin columns.

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- Cells stored in RNAprotect Cell Reagent can also be used in the procedure. Transfer the entire sample, including any material deposited at the bottom of the storage vessel, to a centrifuge tube. Pellet the cells by centrifuging for 5 min at 5000 x *g*, and remove the supernatant by pipetting (if necessary, thaw the sample before centrifuging). Proceed immediately to step 2.
 - Buffer RLT Plus and Buffer RW1 contain a guanidine salt and are therefore not compatible with disinfecting reagents containing bleach. See page 5 for safety information.
 - Perform all steps of the procedure at room temperature (15–25°C). During the procedure, work quickly.
 - Perform all centrifugation steps at 20–25°C in a standard microcentrifuge. Ensure that the centrifuge does not cool below 20°C.

Things to do before starting

- If purifying RNA from cell lines rich in RNases, we recommend adding β -mercaptoethanol (β -ME) to Buffer RLT Plus before use. Add 10 μ l β -ME per 1 ml Buffer RLT Plus. Dispense in a fume hood and wear appropriate protective clothing. Buffer RLT Plus containing β -ME can be stored at room temperature (15–25°C) for up to 1 month. Alternatively, add 20 μ l of 2 M dithiothreitol (DTT) per 1 ml Buffer RLT Plus. The stock solution of 2 M DTT in water should be prepared fresh or frozen in single-use aliquots. Buffer RLT Plus containing DTT can be stored at room temperature for up to 1 month.

- When processing <500 cells, carrier RNA may be added to the lysate before homogenization (see “Carrier RNA”, page 17). Before using for the first time, dissolve the carrier RNA (310 µg) in 1 ml RNase-free water. Store this stock solution at –30°C to –15°C, and use it to make fresh dilutions for each set of RNA preps. The concentration of this stock solution is 310 µg/ml (i.e., 310 ng/µl). To make a working solution (4 ng/µl) for 10 preps, add 5 µl stock solution to 34 µl Buffer RLT Plus and mix by pipetting. Add 6 µl of this diluted solution to 54 µl Buffer RLT Plus to give a working solution of 4 ng/µl. Add 5 µl of this solution to the lysate in step 3. **Do not add the carrier RNA to the lysate if purifying RNA for use in oligo-dT-based amplification.**
- Buffer RPE is supplied as a concentrate. Before using for the first time, add 4 volumes of ethanol (96–100%) as indicated on the bottle to obtain a working solution.
- Before using the kit for the first time, prepare 80% ethanol by mixing 24 ml ethanol (96–100%) and 6 ml RNase-free water (supplied). The procedure also requires 70% ethanol, which can be prepared by diluting ethanol (96–100%) with distilled water (not supplied).
- Buffer RLT Plus may form a precipitate during storage. If necessary, redissolve by warming, and then place at room temperature.

Procedure

1. Harvest cells according to step 1a or 1b.
 - 1a. Cells grown in suspension (do not use more than 5×10^5 cells):

Determine the number of cells. Pellet the appropriate number of cells by centrifuging for 5 min at 300 x g in a centrifuge tube (not supplied). Carefully remove all supernatant by aspiration, and proceed to step 2.

Note: Incomplete removal of cell-culture medium will inhibit lysis and dilute the lysate, affecting the conditions for DNA removal and RNA purification. Both effects may reduce RNA yield and purity.

1b. Cells grown in a monolayer (do not use more than 5×10^5 cells):

Cells grown in a monolayer in cell-culture vessels can be either lysed directly in the vessel (up to 10 cm diameter) or trypsinized and collected as a cell pellet prior to lysis. Cells grown in a monolayer in cell-culture flasks should always be trypsinized.

To lyse cells directly:

Determine the number of cells. Completely aspirate the cell-culture medium, and proceed immediately to step 2.

Note: Incomplete removal of cell-culture medium will inhibit lysis and dilute the lysate, affecting the conditions for DNA removal and RNA purification. Both effects may reduce RNA yield and purity.

To trypsinize and collect cells:

Determine the number of cells. Aspirate the medium, and wash the cells with PBS. Aspirate the PBS, and add 0.10–0.25% trypsin in PBS. After the cells detach from the dish or flask, add medium (containing serum to inactivate the trypsin), transfer the cells to an RNase-free glass or polypropylene centrifuge tube (not supplied), and centrifuge at $300 \times g$ for 5 min. Completely aspirate the supernatant, and proceed to step 2.

Note: Incomplete removal of cell-culture medium will inhibit lysis and dilute the lysate, affecting the conditions for DNA removal and RNA purification. Both effects may reduce RNA yield and purity.

2. Disrupt the cells by adding Buffer RLT Plus.

For pelleted cells, loosen the cell pellet thoroughly by flicking the tube. Add 350 μ l Buffer RLT Plus. Vortex or pipet mix, and proceed to step 3.

If processing $\leq 1 \times 10^5$ cells, 75 μ l Buffer RLT Plus can be added instead. This allows cell pelleting in smaller tubes. Pipet up and down to lyse the cells.

Note: Incomplete loosening of the cell pellet may lead to inefficient lysis and reduced RNA yields.

For direct lysis of cells grown in a monolayer, add 350 μ l Buffer RLT Plus to the cell-culture dish. Collect the lysate with a rubber policeman. Pipet the lysate into a

microcentrifuge tube (not supplied). Vortex or pipet to mix, and ensure that no cell clumps are visible before proceeding to step 3.

If processing $\leq 1 \times 10^5$ cells, 75 μ l Buffer RLT Plus can be added instead. This may be necessary for multiwell plates and cell-culture dishes. Pipet up and down to lyse the cells.

3. Homogenize the lysate according to step 3a, 3b or 3c.

See “Disrupting and homogenizing starting material”, page 15, for more details on homogenization. If processing $\leq 1 \times 10^5$ cells, they can be homogenized by vortexing for 1 min. After homogenization, proceed to step 4.

Note: If only 75 μ l Buffer RLT Plus was used in step 2, transfer the lysate to a new 1.5 ml microcentrifuge tube, and adjust the volume to 350 μ l with Buffer RLT Plus. Vortex for 1 min to homogenize and proceed to step 4.

Note: If processing <500 cells, 20 ng carrier RNA (5 μ l of a 4 ng/ μ l solution) may be added to the lysate before homogenization. Prepare the carrier RNA as described in “Things to do before starting”.

Note: Incomplete homogenization leads to significantly reduced RNA yields and can cause clogging of the gDNA Eliminator and RNeasy MinElute spin columns.

Homogenization with the TissueRuptor or QIAshredder homogenizer generally results in higher RNA yields than with a syringe and needle.

3a. Pipet the lysate directly into a QIAshredder spin column (not supplied) placed in a 2 ml collection tube, and centrifuge for 2 min at full speed. Proceed to step 4.

3b. Place the tip of the TissueRuptor disposable probe into the lysate and operate the TissueRuptor at full speed until the lysate is homogenous (usually 30 s). Proceed to step 4.

Note: To avoid damage to the TissueRuptor II and disposable probe during operation, make sure the tip of the probe remains submerged in the buffer.

3c. Pass the lysate at least 5 times through a blunt 20-gauge needle (0.9 mm diameter) fitted to an RNase-free syringe. Proceed to step 4.

4. Transfer the homogenized lysate to a gDNA Eliminator spin column placed in a 2 ml collection tube (supplied). Centrifuge for 30 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm). Discard the column, and save the flowthrough.

Note: Make sure that no liquid remains on the column membrane after centrifugation. If necessary, repeat the centrifugation until all liquid has passed through the membrane.

The remaining steps of this protocol allow the purification of RNA molecules longer than 200 nucleotides. If purification of total RNA containing small RNAs, such as miRNA is desired, follow steps 1–6 in Appendix D on page 57 instead of steps 5–11 in this protocol.

5. Add 1 volume (usually 350 μ l) of 70% ethanol to the flow-through from step 4, and mix well by pipetting. Do not centrifuge. Proceed immediately to step 6.

Note: The volume of 70% ethanol to add may be less than 350 μ l if some lysate was lost during homogenization and DNA removal.

Note: When purifying RNA from certain cell lines, precipitates may be visible after addition of ethanol. This does not affect the procedure.

6. Transfer the sample, including any precipitate that may have formed, to an RNeasy MinElute spin column placed in a 2 ml collection tube (supplied). Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm). Discard the flow-through.*

Optional: If recovery of protein is desired, keep the flow-through on ice and follow steps 1–5 in Appendix E on page 60.

Reuse the collection tube in step 7.

* Flow-through contains Buffer RLT Plus or Buffer RW1 and is therefore not compatible with bleach. See page 6 for safety information.

-
7. Add 700 μ l Buffer RW1 to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the flow-through.*

Reuse the collection tube in step 8.

Note: After centrifugation, carefully remove the RNeasy MinElute spin column from the collection tube so that the column does not contact the flow-through. Be sure to empty the collection tube completely.

8. Add 500 μ l Buffer RPE to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the flow-through.

Reuse the collection tube in step 9.

Note: Buffer RPE is supplied as a concentrate. Ensure that ethanol is added to Buffer RPE before use (see “Things to do before starting”).

9. Add 500 μ l of 80% ethanol to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 2 min at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the collection tube with the flow-through.

Prepare the 80% ethanol with ethanol (96–100%) and the RNase-free water supplied with the kit.

Note: After centrifugation, carefully remove the RNeasy MinElute spin column from the collection tube so that the column does not contact the flow-through. Otherwise, carryover of ethanol will occur.

10. Place the RNeasy MinElute spin column in a new 2 ml collection tube (supplied). Open the lid of the spin column, and centrifuge at full speed for 5 min. Discard the collection tube with the flow-through.

To avoid damage to their lids, place the spin columns into the centrifuge with at least one empty position between columns. Orient the lids so that they point in a direction opposite to the rotation of the rotor (e.g., if the rotor rotates clockwise, orient the lids counterclockwise).

It is important to dry the spin column membrane, since residual ethanol may interfere with downstream reactions. Centrifugation with the lids open ensures that no ethanol is carried over during RNA elution.

11. Place the RNeasy MinElute spin column in a new 1.5 ml collection tube (supplied). Add 14 μ l RNase-free water directly to the center of the spin column membrane. Close the lid gently, and centrifuge for 1 min at full speed to elute the RNA.

As little as 10 μ l RNase-free water can be used for elution if a higher RNA concentration is required, but the yield will be reduced by approximately 20%. Do not elute with less than 10 μ l RNase-free water, as the spin column membrane will not be sufficiently hydrated.

The dead volume of the RNeasy MinElute spin column is 2 μ l: elution with 14 μ l RNase-free water results in a 12 μ l eluate.

For RT-PCR and real-time RT-PCR with the purified RNA, QIAGEN offers a range of optimized, ready-to-use kits that provide highly specific and sensitive results. For details, visit www.qiagen.com/PCR. For whole transcriptome amplification (WTA) of limited amounts of RNA, we recommend the QuantiTect Whole Transcriptome Kit.

Protocol: Purification of Total RNA from Animal and Human Tissues

This protocol is for the purification of total RNA from easy-to-lyse animal and human tissues. For total RNA purification from frozen, microdissected tissue samples, see page 39.

Determining the correct amount of starting material

It is essential to use the correct amount of starting material to obtain optimal RNA yield and purity. A maximum amount of 5 mg fresh or frozen tissue or 2–3 mg RNAprotect- or Allprotect-stabilized tissue (which is partially dehydrated) can generally be processed. For most tissues, the DNA removal capacity of the gDNA Eliminator spin column, the RNA binding capacity of the RNeasy MinElute spin column and the lysing capacity of Buffer RLT Plus will not be exceeded by these amounts. Typical RNA yields from various tissues are given in Table 2 (page 13).

For maximum RNA yields from liver, 50% ethanol (instead of 70% ethanol) should be used in step 4 of the procedure.

Some tissues, such as spleen, parts of brain, lung and thymus, tend to form precipitates during the procedure. However, this does not affect RNA purification.

Do not overload the gDNA Eliminator spin column, as this will lead to copurification of DNA with RNA. Do not overload the RNeasy MinElute spin column, as this will significantly reduce RNA yield and quality.

Weighing tissue is the most accurate way to quantitate the amount of starting material. As a guide, a 1.5 mm cube (3.4 mm³) of most animal tissues weighs 3.5–4.5 mg.

Important points before starting

- If using the RNeasy Plus Micro Kit for the first time, read “Important Notes” (page 11).
- If preparing RNA for the first time, read Appendix A (page 49).
- If using the TissueRuptor II, ensure that you are familiar with operating it by referring to the *TissueRuptor II User Manual* and *TissueRuptor II Handbook*.
- If using the TissueLyser, ensure that you are familiar with operating it by referring to the operating instructions and *TissueLyser II Handbook*.
- For optimal results, stabilize harvested tissues immediately in RNAprotect Tissue Reagent (see the *RNAprotect Handbook*) or Allprotect Tissue Reagent (see the *Allprotect Tissue Reagent Handbook*). Tissues can be stored in the reagent at 37°C for up to 1 day, at 15–25°C for up to 7 days, or at 2–8°C for up to 4 weeks (RNAprotect) or 6 months (Allprotect). Alternatively, tissues can be archived at –30°C to –15°C or –90°C to –65°C.
- Fresh, frozen or RNAprotect- or Allprotect-stabilized tissue can be used. Tissues can be stored at –90°C to –65°C for several months. Flash-freeze tissues in liquid nitrogen, and immediately transfer to –90°C to –65°C. Do not allow tissues to thaw during weighing or handling prior to disruption in Buffer RLT Plus. Homogenized tissue lysates from step 2 can also be stored at –90°C to –65°C for several months. Incubate frozen lysates at 37°C in a water bath until completely thawed and salts are dissolved before continuing with step 3. Avoid prolonged incubation, which may compromise RNA integrity.
- Buffer RLT Plus and Buffer RW1 contain a guanidine salt and are therefore not compatible with disinfecting reagents containing bleach. See page 5 for safety information.
- Perform all steps of the procedure at room temperature (15–25°C). During the procedure, work quickly.
- Perform all centrifugation steps at 20–25°C in a standard microcentrifuge. Ensure that the centrifuge does not cool below 20°C.

Things to do before starting

- β -Mercaptoethanol (β -ME) must be added to Buffer RLT Plus before use. Add 10 μ l β -ME per 1 ml Buffer RLT Plus. Dispense in a fume hood and wear appropriate protective clothing. Buffer RLT Plus containing β -ME can be stored at room temperature (15–25°C) for up to 1 month. Alternatively, add 20 μ l of 2 M dithiothreitol (DTT) per 1 ml Buffer RLT Plus. The stock solution of 2 M DTT in water should be prepared fresh or frozen in single-use aliquots. Buffer RLT Plus containing DTT can be stored at room temperature for up to 1 month.
- When processing less than about 2 μ g tissue, carrier RNA may be added to the lysate before homogenization (see “Carrier RNA”, page 17). Before using for the first time, dissolve the carrier RNA (310 μ g) in 1 ml RNase-free water. Store this stock solution at –30°C to –15°C, and use it to make fresh dilutions for each set of RNA preps. The concentration of this stock solution is 310 μ g/ml (i.e., 310 ng/ μ l). To make a working solution (4 ng/ μ l) for 10 preps, add 5 μ l stock solution to 34 μ l Buffer RLT Plus and mix by pipetting. Add 6 μ l of this diluted solution to 54 μ l Buffer RLT Plus to give a working solution of 4 ng/ μ l. Add 5 μ l of this solution to the lysate in step 2. **Do not add the carrier RNA to the lysate if purifying RNA for use in oligo-dT-based amplification.**
- Buffer RPE is supplied as a concentrate. Before using for the first time, add 4 volumes of ethanol (96–100%) as indicated on the bottle to obtain a working solution.
- Before using the kit for the first time, prepare 80% ethanol by mixing 24 ml ethanol (96–100%) and 6 ml RNase-free water (supplied). The procedure also requires 70% ethanol, which can be prepared by diluting ethanol (96–100%) with distilled water (not supplied).
- Buffer RLT Plus may form a precipitate during storage. If necessary, redissolve by warming and then place at room temperature.

Procedure

- Excise the tissue sample from the animal or remove it from storage. Determine the amount of tissue. Do not use more than 5 mg. Proceed immediately to step 2.

Weighing tissue is the most accurate way to determine the amount. If necessary, cut the tissue on a clean surface and weigh the piece to be used.

For RNAprotect- or Allprotect-stabilized tissues: Remove the tissue from the stabilization reagent using forceps and be sure to remove any crystals that may have formed. RNA in RNAprotect- or Allprotect-stabilized tissues is protected during cutting and weighing of tissues at ambient temperature (15–25°C). It is not necessary to cut the tissues on ice or dry ice or in a refrigerated room. Remaining tissues can be stored in RNAprotect Tissue Reagent or Allprotect Reagent. Previously stabilized tissues can be stored at –90°C to –65°C without the reagent.

For unstabilized fresh or frozen tissues: RNA in harvested tissues is not protected until the tissues are treated with RNAprotect Tissue Reagent or Allprotect Reagent, flash-frozen or disrupted and homogenized in step 2. Frozen tissues should not be allowed to thaw during handling. The relevant procedures should be carried out as quickly as possible. Remaining fresh tissues can be placed into RNAprotect Tissue Reagent to stabilize RNA or in Allprotect Tissue Reagent to stabilize DNA, RNA and protein. However, previously frozen tissues thaw too slowly in the reagent, preventing the reagent from diffusing into the tissues quickly enough to prevent RNA degradation.

12. Disrupt the tissue and homogenize the lysate in Buffer RLT Plus (do not use more than 5 mg tissue) according to step 2a, 2b or 2c.

See “Disrupting and homogenizing starting material”, page 15, for more details on disruption and homogenization.

Note: Ensure that β -ME (or DTT) is added to Buffer RLT Plus before use (see “Things to do before starting”).

Note: If processing <2 μ g tissue, 20 ng carrier RNA (5 μ l of a 4 ng/ μ l solution) may be added to the lysate before homogenization. Prepare the carrier RNA as described in “Things to do before starting”.

After storage in RNAprotect Tissue Reagent or Allprotect Reagent, tissues may become slightly harder than fresh or thawed tissues. Disruption and homogenization using standard methods is usually not a problem.

Note: Incomplete homogenization leads to significantly reduced RNA yields and can cause clogging of the gDNA Eliminator and RNeasy MinElute spin columns. Homogenization with the TissueRuptor or TissueLyser II generally results in higher RNA yields than with other methods.

12a. Disruption and homogenization using the TissueRuptor II:

- Place the tissue in a suitably sized vessel. Add 350 µl Buffer RLT Plus.

Note: Use a suitably sized vessel with sufficient extra headspace to accommodate foaming, which may occur during homogenization.

Generally, round-bottomed tubes allow more efficient disruption and homogenization than conical-bottomed tubes.

- Place the tip of the disposable probe into the vessel and operate the TissueRuptor II at full speed until the lysate is homogeneous (usually 30 s). Proceed to step 3.

Note: To avoid damage to the TissueRuptor II and disposable probe during operation, make sure the tip of the probe remains submerged in the buffer.

Foaming may occur during homogenization. If this happens, let the homogenate stand at room temperature (15–25°C) for 2–3 min until the foam subsides before continuing with the procedure.

12b. Disruption and homogenization using the TissueLyser II:

- Place the tissues in 2 ml microcentrifuge tubes containing one stainless steel bead (5 mm mean diameter).

If handling fresh or frozen tissue samples, keep the tubes on dry ice.

- Place the tubes at room temperature. Immediately add 350 µl Buffer RLT Plus per tube.
- Place the tubes in the TissueLyser Adapter Set 2 x 24.

- Operate the TissueLyser II for 2 min at 20 Hz.
The time depends on the tissue being processed and can be extended until the tissue is completely homogenized.
- Rearrange the collection tubes so that the outermost tubes are innermost and the innermost tubes are outermost. Operate the TissueLyser II for another 2 min at 20 Hz. Rearranging the tubes allows even homogenization.
- Carefully pipet the lysates into new microcentrifuge tubes (not supplied). Proceed to step 3.
Do not reuse the stainless steel beads.

12c. Disruption using a mortar and pestle followed by homogenization using a QIAshredder homogenizer or a needle and syringe:

- Immediately place the weighed tissue in liquid nitrogen, and grind thoroughly with a mortar and pestle.
- Decant tissue powder and liquid nitrogen into an RNase-free, liquid-nitrogen-cooled, 2 ml microcentrifuge tube (not supplied). Allow the liquid nitrogen to evaporate, but do not allow the tissue to thaw.
- Add 350 μ l Buffer RLT Plus.
- Pipet the lysate directly into a QIAshredder spin column placed in a 2 ml collection tube, and centrifuge for 2 min at full speed. Alternatively, pass the lysate at least 5 times through a blunt 20-gauge needle fitted to an RNase-free syringe. Proceed to step 3.

13. Centrifuge the lysate for 3 min at full speed. Carefully remove the supernatant by pipetting, and transfer it to a gDNA Eliminator spin column placed in a 2 ml collection tube (supplied). Centrifuge for 30 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm). Discard the column, and save the flow-through.

This step is important, as it removes insoluble material that could clog the gDNA Eliminator spin column and interfere with DNA removal. In some preparations, very small amounts of insoluble material will be present after the 3-min centrifugation, making the pellet invisible.

Note: Make sure that no liquid remains on the column membrane after centrifugation. If necessary, repeat the centrifugation until all liquid has passed through the membrane.

14. Add 1 volume (usually 350 μ l) of 70% ethanol to the flow-through from step 3, and mix well by pipetting. Do not centrifuge. Proceed immediately to step 5.

Note: The volume of 70% ethanol to add may be less than 350 μ l if some lysate was lost during homogenization and DNA removal.

Note: Precipitates may be visible after addition of ethanol, but this does not affect the procedure.

Note: For maximum RNA yields from liver, use 50% ethanol instead of 70% ethanol.

15. Transfer the sample, including any precipitate that may have formed, to an RNeasy MinElute spin column placed in a 2 ml collection tube (supplied). Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm). Discard the flowthrough.*

Optional: If recovery of protein is desired, keep the flow-through on ice and follow steps 1–5 in Appendix E on page 60.

Reuse the collection tube in step 6.

* Flow-through contains Buffer RLT Plus or Buffer RW1 and is therefore not compatible with bleach. See page 6 for safety information.

16. Add 700 μ l Buffer RW1 to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the flow-through.*

Reuse the collection tube in step 7.

Note: After centrifugation, carefully remove the RNeasy MinElute spin column from the collection tube so that the column does not contact the flow-through. Be sure to empty the collection tube completely.

17. Add 500 μ l Buffer RPE to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the flow-through.

Reuse the collection tube in step 8.

Note: Buffer RPE is supplied as a concentrate. Ensure that ethanol is added to Buffer RPE before use (see “Things to do before starting”).

18. Add 500 μ l of 80% ethanol to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 2 min at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the collection tube with the flow-through.

Prepare the 80% ethanol with ethanol (96–100%) and the RNase-free water supplied with the kit.

Note: After centrifugation, carefully remove the RNeasy MinElute spin column from the collection tube so that the column does not contact the flow-through. Otherwise, carryover of ethanol will occur.

19. Place the RNeasy MinElute spin column in a new 2 ml collection tube (supplied). Open the lid of the spin column, and centrifuge at full speed for 5 min. Discard the collection tube with the flow-through.

To avoid damage to their lids, place the spin columns into the centrifuge with at least one empty position between columns. Orient the lids so that they point in a direction opposite to the rotation of the rotor (e.g., if the rotor rotates clockwise, orient the lids counterclockwise).

It is important to dry the spin column membrane, since residual ethanol may interfere with downstream reactions. Centrifugation with the lids open ensures that no ethanol is carried over during RNA elution.

20. Place the RNeasy MinElute spin column in a new 1.5 ml collection tube (supplied). Add 14 μ l RNase-free water directly to the center of the spin column membrane. Close the lid gently, and centrifuge for 1 min at full speed to elute the RNA.

As little as 10 μ l RNase-free water can be used for elution if a higher RNA concentration is required, but the yield will be reduced by approximately 20%. Do not elute with less than 10 μ l RNase-free water, as the spin column membrane will not be sufficiently hydrated.

The dead volume of the RNeasy MinElute spin column is 2 μ l: elution with 14 μ l RNase-free water results in a 12 μ l eluate.

For RT-PCR and real-time RT-PCR with the purified RNA, QIAGEN offers a range of optimized, ready-to-use kits that provide highly specific and sensitive results. For details, visit www.qiagen.com/PCR. For whole transcriptome amplification (WTA) of limited amounts of RNA, we recommend the QuantiTect Whole Transcriptome Kit.

Protocol: Purification of Total RNA from Microdissected Cryosections

This protocol is for the purification of total RNA from frozen, microdissected samples of animal and human tissues. For total RNA purification from microdissected, formalin-fixed samples, we recommend the RNeasy FFPE Kit (cat. no. 74404).

Laser-microdissected tissue specimens present a particular challenge for molecular analysis, as nucleic acids must be purified from very small amounts of starting material. In addition, fixation and staining steps may compromise the integrity of RNA, and it may be necessary either to modify fixation protocols or to use cryosections from flash-frozen specimens to minimize this problem.

A wide range of equipment and consumables for sectioning, staining and microdissection of specimens is available from Leica (www.leica-microsystems.com) and P.A.L.M. Microlaser Technologies (www.palm-mikrolaser.com).

Important points before starting

- If using the RNeasy Plus Micro Kit for the first time, read “Important Notes” (page 11).
- If preparing RNA for the first time, read Appendix A (page 49).
- To minimize RNA degradation, avoid prolonged storage of unstabilized samples at room temperature (15–25°C). RNA in tissues is not protected before flash-freezing in liquid nitrogen.
- Tissue lysates from step 3 can be stored at –90°C to –65°C for several months. Incubate frozen lysates at 37°C in a water bath until completely thawed and salts are dissolved before continuing with step 4. Avoid prolonged incubation, which may compromise RNA integrity.

- Buffer RLT Plus and Buffer RW1 contain a guanidine salt and are therefore not compatible with disinfecting reagents containing bleach. See page 5 for safety information.
- Perform all steps of the procedure at room temperature (15–25°C). During the procedure, work quickly.
- Perform all centrifugation steps at 20–25°C in a standard microcentrifuge. Ensure that the centrifuge does not cool below 20°C.

Things to do before starting

- β -Mercaptoethanol (β -ME) must be added to Buffer RLT Plus before use. Add 10 μ l β -ME per 1 ml Buffer RLT Plus. Dispense in a fume hood and wear appropriate protective clothing. Buffer RLT Plus containing β -ME can be stored at room temperature (15–25°C) for up to 1 month. Alternatively, add 20 μ l of 2 M dithiothreitol (DTT) per 1 ml Buffer RLT Plus. The stock solution of 2 M DTT in water should be prepared fresh or frozen in single-use aliquots. Buffer RLT Plus containing DTT can be stored at room temperature for up to 1 month.
- When processing <500 cells, carrier RNA may be added to the lysate before homogenization (see “Carrier RNA”, page 17). Before using for the first time, dissolve the carrier RNA (310 μ g) in 1 ml RNase-free water. Store this stock solution at –30°C to –15°C, and use it to make fresh dilutions for each set of RNA preps. The concentration of this stock solution is 310 μ g/ml (i.e., 310 ng/ μ l). To make a working solution (4 ng/ μ l) for 10 preps, add 5 μ l stock solution to 34 μ l Buffer RLT Plus and mix by pipetting. Add 6 μ l of this diluted solution to 54 μ l Buffer RLT Plus to give a working solution of 4 ng/ μ l. Add 5 μ l of this solution to the lysate in step 2. **Do not add the carrier RNA to the lysate if purifying RNA for use in oligo-dT-based amplification.**
- Buffer RPE is supplied as a concentrate. Before using for the first time, add 4 volumes of ethanol (96–100%) as indicated on the bottle to obtain a working solution.

- Before using the kit for the first time, prepare 80% ethanol by mixing 24 ml ethanol (96–100%) and 6 ml RNase-free water (supplied). The procedure also requires 70% ethanol, which can be prepared by diluting ethanol (96–100%) with distilled water (not supplied).
- Buffer RLT Plus may form a precipitate during storage. If necessary, redissolve by warming, and then place at room temperature.

Procedure

1. Collect the sample directly into an appropriate volume of Buffer RLT Plus (the volume depends on the collection vessel used for microdissection, but should not be greater than 65 μ l [Leica® instruments] or 300 μ l [other instruments]).

Note: Ensure that β -ME (or DTT) is added to Buffer RLT Plus before use (see “Things to do before starting”).

2. If necessary, transfer the sample and buffer to a larger vessel (e.g., 1.5 or 2 ml tube). Adjust the volume to 350 μ l with Buffer RLT Plus.

Note: If processing <500 cells, 20 ng carrier RNA (5 μ l of a 4 ng/ μ l solution) may be added to the lysate before homogenization. Prepare the carrier RNA as described in “Things to do before starting”.

3. Vortex the sample for 30 s.

No further homogenization is necessary.

4. Transfer the sample to a gDNA Eliminator spin column placed in a 2 ml collection tube (supplied). Centrifuge for 30 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm). Discard the column, and save the flow-through.

Note: Make sure that no liquid remains on the column membrane after centrifugation. If necessary, repeat the centrifugation until all liquid has passed through the membrane.

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5. Add 1 volume (usually 350 μ l) of 70% ethanol to the flow-through from step 4, and mix well by pipetting. Do not centrifuge. Proceed immediately to step 6.

Note: The volume of 70% ethanol to add may be less than 350 μ l if some lysate was lost during homogenization and DNA removal.

Note: Precipitates may be visible after addition of ethanol. This does not affect the procedure.

6. Transfer the sample, including any precipitate that may have formed, to an RNeasy MinElute spin column placed in a 2 ml collection tube (supplied). Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm). Discard the flow-through.*

Reuse the collection tube in step 7.

7. Add 700 μ l Buffer RW1 to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the flow-through.*

Reuse the collection tube in step 8.

Note: After centrifugation, carefully remove the RNeasy MinElute spin column from the collection tube so that the column does not contact the flow-through. Be sure to empty the collection tube completely.

8. Add 500 μ l Buffer RPE to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the flow-through.

Reuse the collection tube in step 9.

Note: Buffer RPE is supplied as a concentrate. Ensure that ethanol is added to Buffer RPE before use (see “Things to do before starting”).

* Flow-through contains Buffer RLT or Buffer RW1 and is therefore not compatible with bleach. See page 6 for safety information.

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9. Add 500 μ l of 80% ethanol to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 2 min at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the flow-through and collection tube.

Prepare the 80% ethanol with ethanol (96–100%) and the RNase-free water supplied with the kit.

Note: After centrifugation, carefully remove the RNeasy MinElute spin column from the collection tube so that the column does not contact the flow-through. Otherwise, carryover of ethanol will occur.

10. Place the RNeasy MinElute spin column in a new 2 ml collection tube (supplied). Open the lid of the spin column, and centrifuge at full speed for 5 min. Discard the collection tube with the flow-through.

To avoid damage to their lids, place the spin columns into the centrifuge with at least one empty position between columns. Orient the lids so that they point in a direction opposite to the rotation of the rotor (e.g., if the rotor rotates clockwise, orient the lids counterclockwise).

It is important to dry the spin column membrane, since residual ethanol may interfere with downstream reactions. Centrifugation with the lids open ensures that no ethanol is carried over during RNA elution.

11. Place the RNeasy MinElute spin column in a new 1.5 ml collection tube (supplied). Add 14 μ l RNase-free water directly to the center of the spin column membrane. Close the lid gently, and centrifuge for 1 min at full speed to elute the RNA.

As little as 10 μ l RNase-free water can be used for elution if a higher RNA concentration is required, but the yield will be reduced by approximately 20%. Do not elute with less than 10 μ l RNase-free water, as the spin column membrane will not be sufficiently hydrated.

The dead volume of the RNeasy MinElute spin column is 2 μ l: elution with 14 μ l RNase-free water results in a 12 μ l eluate.

For RT-PCR and real-time RT-PCR with the purified RNA, QIAGEN offers a range of optimized, ready-to-use kits that provide highly specific and sensitive results. For details, visit www.qiagen.com/PCR. For whole transcriptome amplification (WTA) of limited amounts of RNA, we recommend the QuantiTect Whole Transcriptome Kit.

Troubleshooting Guide

This troubleshooting guide may be helpful in solving any problems that may arise. The scientists in QIAGEN Technical Services are always happy to answer any questions you may have about either the information and protocols in this handbook or molecular biology applications (for contact information, see back cover or visit www.qiagen.com).

Comments and suggestions

Clogged RNeasy MinElute spin column

- | | |
|---|--|
| a) Inefficient disruption and/or homogenization | See “Disrupting and homogenizing starting materials” (page 15) for details on disruption and homogenization methods.

Increase <i>g</i> -force and centrifugation time if necessary.

In subsequent preparations, reduce the amount of starting material (see the individual protocols) and/or increase the homogenization time. |
| b) Too much starting material | Reduce the amount of starting material (see the individual protocols). It is essential to use the correct amount of starting material. |
| c) Centrifugation temperature too low | The centrifugation temperature should be 20–25°C. Some centrifuges may cool to below 20°C even when set at 20°C. This can cause formation of precipitates that can clog the spin column. If this happens, set the centrifugation temperature to 25°C. Warm the lysate to 37°C before transferring it to the gDNA Eliminator spin column. |

Low RNA yield

- | | |
|---|---|
| a) Insufficient disruption and homogenization | See “Disrupting and homogenizing starting materials” (page 15) for details on disruption and homogenization methods.

In subsequent preparations, reduce the amount of starting material (see the individual protocols) and/or increase the volume of lysis buffer and the homogenization time. |
|---|---|

Comments and suggestions

- | | | |
|----|--|---|
| b) | Too much starting material | In subsequent preparations, reduce the amount of starting material (see the individual protocols). It is essential to use the correct amount of starting material. |
| c) | RNA still bound to spin column membrane | Repeat RNA elution, but incubate the RNeasy MinElute spin column on the benchtop for 10 min with RNase-free water before centrifuging. |
| d) | Ethanol carryover | <p>After the wash with 80% ethanol, be sure to centrifuge at full speed for 5 min to dry the RNeasy MinElute spin column membrane.</p> <p>After centrifugation, carefully remove the RNeasy MinElute spin column from the collection tube so that the column does not contact the flow-through. Otherwise, carryover of ethanol will occur.</p> |
| e) | Incomplete removal of cell-culture medium (cell samples) | When processing cultured cells, ensure complete removal of cell-culture medium after harvesting cells (see protocol, page 21). |
| f) | 80% ethanol not made with RNase-free water | The 80% ethanol used to wash the RNeasy MinElute spin column membrane must be free of RNases. Be sure to prepare the 80% ethanol using ethanol (96–100%) and the RNase-free water supplied with the kit, as described in “Things to do before starting” in each protocol. |

Low or no recovery of RNA

- | | | |
|----|--|---|
| a) | RNase-free water incorrectly dispensed | Pipet RNase-free water to the center of the RNeasy MinElute spin column membrane to ensure that the membrane is completely covered. |
| b) | Ethanol carryover | <p>After the wash with 80% ethanol, be sure to centrifuge at full speed for 5 min to dry the RNeasy MinElute spin column membrane.</p> <p>After centrifugation, carefully remove the RNeasy MinElute spin column from the collection tube so that the column does not contact the flow-through. Otherwise, carryover of ethanol will occur.</p> |

Comments and suggestions

Low A_{260}/A_{280} value

Water use to dilute RNA for A_{260}/A_{280} measurement

Use 10 mM Tris-Cl, pH 7.5, not RNase-free water, to dilute the sample before measuring purity (see Appendix B, page **Error! Bookmark not defined.**)

RNA degraded

- a) Inappropriate handling of starting material
- Ensure that tissue samples are properly stabilized and stored in RNAprotect Tissue Reagent or Allprotect Tissue Reagent. For frozen cell pellets or frozen tissue samples, ensure that they were flash-frozen immediately in liquid nitrogen and properly stored at -90°C to -65°C . Perform the RNeasy procedure quickly, especially the first few steps. See Appendix A (page 49) and "Handling and storing starting material" (page 13).
- b) RNase contamination
- Although all RNeasy buffers have been tested and are guaranteed RNase-free, RNases can be introduced during use. Be certain not to introduce any RNases during the RNeasy procedure or later handling. See Appendix A (page 49) for general remarks on handling RNA. Do not put RNA samples into a vacuum dryer or microcentrifuge that has been used in DNA preparation where RNases may have been used.
- c) 80% ethanol not made with RNase-free water
- The 80% ethanol used to wash the RNeasy MinElute spin column membrane must be free of RNases. Be sure to prepare the 80% ethanol using ethanol (96–100%) and the RNase-free water supplied with the kit, as described in "Things to do before starting" in each protocol.

Contamination of RNA with DNA affects downstream applications

- a) Cell number too high
- For some cell types, the efficiency of DNA binding to the gDNA Eliminator spin column may be reduced when processing very high cell numbers. If the eluted RNA contains substantial DNA contamination, try processing smaller cell numbers.

Comments and suggestions

- | | | |
|----|--|--|
| b) | Incomplete removal of cell-culture medium or stabilization reagent | Be sure to remove any excess cell-culture medium or stabilization reagent to prevent significant dilution of the lysis buffer. The gDNA Eliminator spin column will not bind DNA effectively if the lysis buffer is diluted. |
| c) | Tissue has high DNA content | For certain tissues with extremely high DNA content (e.g., thymus), trace amounts of DNA may pass through the gDNA Eliminator spin column. Try using smaller samples. |

RNA does not perform well in downstream experiments

- | | | |
|----|---|---|
| a) | Ethanol carryover | After the wash with 80% ethanol, be sure to centrifuge at full speed for 5 min to dry the RNeasy MinElute spin column membrane.

After centrifugation, carefully remove the RNeasy MinElute spin column from the collection tube so that the column does not contact the flow-through. Otherwise, carryover of ethanol will occur. |
| b) | Salt carryover during elution | Ensure that buffers are at 20–30°C.

Ensure that the correct buffer is used for each step of the procedure.

When reusing collection tubes between washing steps, remove residual flow-through from the rim by blotting on clean paper towels. |
| c) | Reverse transcription with too small an amount of RNA | When performing reverse transcription with very small amounts of RNA, we recommend using the Sensiscript® RT Kit, which is specially designed for cDNA synthesis from <50 ng RNA. If synthesizing cDNA for use in real-time PCR, we recommend the QuantiNova® Reverse Transcription Kit, which is compatible with a wide range of RNA amounts (10 pg to 1 µg). For ordering information, see page 62. |

Appendix A: General Remarks on Handling RNA

Handling RNA

Ribonucleases (RNases) are very stable and active enzymes that generally do not require cofactors to function. Since RNases are difficult to inactivate and even minute amounts are sufficient to destroy RNA, do not use any plasticware or glassware without first eliminating possible RNase contamination. Great care should be taken to avoid inadvertently introducing RNases into the RNA sample during or after the purification procedure. To create and maintain an RNase-free environment, the following precautions must be taken during pretreatment and use of disposable and nondisposable vessels and solutions while working with RNA.

General handling

Proper microbiological, aseptic technique should always be used when working with RNA. Hands and dust particles may carry bacteria and molds and are the most common sources of RNase contamination. Always wear latex or vinyl gloves while handling reagents and RNA samples to prevent RNase contamination from the surface of the skin or from dusty laboratory equipment. Change gloves frequently and keep tubes closed whenever possible. Keep purified RNA on ice when aliquots are pipetted for downstream applications. To remove RNase contamination from bench surfaces, nondisposable plasticware and laboratory equipment (e.g., pipets and electrophoresis tanks), use general laboratory reagents. To decontaminate plasticware, rinse with 0.1 M NaOH, 1 mM EDTA,* followed by RNase-free water (see "Solutions", page 50), or rinse with chloroform* if the plasticware is chloroform-resistant. To decontaminate electrophoresis tanks, clean with detergent (e.g., 0.5% SDS),* rinse with RNase-free water, then rinse with ethanol (if the tanks are ethanol resistant) and allow to dry.

* When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, consult the appropriate safety data sheets (SDSs), available from the product supplier.

Disposable plasticware

The use of sterile, disposable polypropylene tubes is recommended throughout the procedure. These tubes are generally RNase-free and do not require pretreatment to inactivate RNases.* When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, consult the appropriate safety data sheets (SDSs), available from the product supplier. Glassware should be treated before use to ensure that it is RNase-free. Glassware used for RNA work should be cleaned with a detergent,* thoroughly rinsed and oven baked at 240°C for at least 4 hours (overnight, if more convenient) before use. Autoclaving alone will not fully inactivate many RNases. Alternatively, glassware can be treated with diethyl pyrocarbonate (DEPC)*, as described in “Solutions” below.

Solutions

Solutions (water and other solutions) should be treated with 0.1% DEPC. DEPC is a strong, but not absolute, inhibitor of RNases. It is commonly used at a concentration of 0.1% to inactivate RNases on glass or plasticware or to create RNase-free solutions and water. DEPC inactivates RNases by covalent modification. Add 0.1 ml DEPC to 100 ml of the solution to be treated and shake vigorously to bring the DEPC into solution. Let the solution incubate for 12 hours at 37°C. Autoclave for 15 minutes to remove any trace of DEPC. DEPC will react with primary amines and cannot be used directly to treat Tris* buffers. DEPC is highly unstable in the presence of Tris buffers and decomposes rapidly into ethanol and CO₂. When preparing Tris buffers, treat water with DEPC first, and then dissolve Tris to make the appropriate buffer. Trace amounts of DEPC will modify purine residues in RNA by carbethoxylation. Carbethoxylated RNA is translated with very low efficiency in cell-free systems. However, its ability to form DNA:RNA or RNA:RNA hybrids is not seriously affected unless a large fraction of the purine residues has been modified. Residual DEPC must always be eliminated from solutions or vessels by autoclaving or heating to 100°C for 15 minutes.

* When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, consult the appropriate safety data sheets (SDSs), available from the product supplier.

Note: RNeasy buffers are guaranteed RNase-free without using DEPC treatment and are therefore free of any DEPC contamination.

Appendix B: Storage, Quantification and Determination of Quality of RNA

Storage of RNA

Purified RNA may be stored at -70°C to -15°C in RNase-free water. Under these conditions, no degradation of RNA is detectable after 1 year.

Quantification of RNA

The concentration of RNA should be determined by measuring the absorbance at 260 nm (A_{260}) in a spectrophotometer (see “Spectrophotometric quantification of RNA” below). For small amounts of RNA, however, it may be difficult to determine amounts photometrically. Small amounts of RNA can be quantified using quantitative RT-PCR or fluorometric quantification.

Spectrophotometric quantification of RNA

Using the QIAxpert UV/VIS Spectrophotometer for microvolume analysis

To determine the concentration of your RNA sample purified with RNeasy QIAGEN kit, use the corresponding RNeasy App on the QIAxpert. For more information, see the QIAxpert product page (www.qiagen.com/qiaxpert-system).

Using a standard spectrophotometer

To ensure significance, A_{260} readings should be greater than 0.15. An absorbance of 1 unit at 260 nm corresponds to 44 μg of RNA per ml ($A_{260} = 1 \rightarrow 4 \mu\text{g}/\text{ml}$). This relation is valid only for measurements at a neutral pH. Therefore, if it is necessary to dilute the RNA sample,

this should be done in a buffer with neutral pH.* As discussed below (see "Purity of RNA", page 53), the ratio between the absorbance values at 260 and 280 nm gives an estimate of RNA purity. When measuring RNA samples, be certain that cuvettes are RNase-free, especially if the RNA is to be recovered after spectrophotometry. This can be accomplished by washing cuvettes with 0.1 M NaOH, 1 mM EDTA,* followed by washing with RNase-free water (see "Solutions", page 50). Use the buffer in which the RNA is diluted to zero the spectrophotometer. An example of the calculation involved in RNA quantification is shown below:

Volume of RNA sample = 100 μ l
Dilution = 10 μ l of RNA sample + 490 μ l of 10 mM Tris-Cl,* pH 7.0
(1/50 dilution)

Measure absorbance of diluted sample in a 1 ml cuvette (RNase-free)

A_{260} = 0.2
Concentration of RNA sample = 44 μ g/ml $\times A_{260}$ \times dilution factor
= 44 μ g/ml \times 0.2 \times 50
= 440 μ g/ml

Total amount = concentration \times volume in milliliters
= 440 μ g/ml \times 0.1 ml
= 44 μ g of RNA

Purity of RNA

The assessment of RNA purity will be performed routinely, when using the QIAxpert with the corresponding RNeasy App. See the QIAxpert user manual for more information (www.qiagen.com/qiexpert-system/user manual)

* When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, consult the appropriate safety data sheets (SDSs), available from the product supplier.

For standard photometric measurements, the ratio of the readings at 260 nm and 280 nm (A_{260}/A_{280}) provides an estimate of the purity of RNA with respect to contaminants, such as protein, that absorb in the UV spectrum. However, the A_{260}/A_{280} ratio is influenced considerably by pH. Since water is not buffered, the pH and the resulting A_{260}/A_{280} ratio can vary greatly when using pure water. Lower pH results in a lower A_{260}/A_{280} ratio and reduced sensitivity to protein contamination.* For accurate values, we recommend measuring absorbance in 10 mM Tris-Cl, pH 7.5. Pure RNA has an A_{260}/A_{280} ratio of 1.9–2.1[†] in 10 mM Tris-Cl, pH 7.5. Always be sure to calibrate the spectrophotometer with the same solution used for dilution. For determination of RNA concentration, however, we recommend dilution of the sample in a buffer with neutral pH since the relationship between absorbance and concentration (A_{260} reading of 1 = 44 $\mu\text{g}/\text{ml}$ RNA) is based on an extinction coefficient calculated for RNA at neutral pH (see “Spectrophotometric quantification of RNA”, page 52).

DNA contamination

No currently available purification method can guarantee that RNA is completely free of DNA, even when it is not visible on an agarose gel. RNeasy Kits will, however, remove the vast majority of cellular DNA. gDNA Eliminator Solution helps to further reduce genomic DNA contamination; however, trace amounts of genomic DNA may still remain, depending on the amount and nature of the sample. For analysis of very low abundance targets, any interference by residual DNA contamination can be detected by performing real-time RT-PCR control experiments in which no reverse transcriptase is added prior to the PCR step.

To prevent any interference by DNA in real-time RT-PCR applications, such as with Applied Biosystems® and Rotor-Gene® instruments, we recommend designing primers that anneal at intron splice junctions so that genomic DNA will not be amplified. QuantiTect Primer Assays from QIAGEN are designed for SYBR® Green-based real-time RT-PCR analysis of RNA sequences (without detection of genomic DNA) where possible (see www.qiagen.com/GeneGlobe). For real-time RT-PCR assays where amplification of genomic

* Wilfinger, W.W., Mackey, M., and Chomczynski, P. (1997) Effect of pH and ionic strength on the spectrophotometric assessment of nucleic acid purity. *BioTechniques* **22**, 474.

[†] Values up to 2.3 are routinely obtained for pure RNA (in 10 mM Tris-Cl, pH 7.5) with some spectrophotometers.

DNA cannot be avoided, we recommend using the QuantiTect Reverse Transcription Kit for reverse transcription. The kit integrates fast cDNA synthesis with rapid removal of genomic DNA contamination (see ordering information, page 62).

Integrity of RNA

The integrity and size distribution of total RNA purified with RNeasy Plus Universal Kits can be checked by denaturing agarose gel electrophoresis and ethidium bromide staining* or by using the QIAxcel system or Agilent® 2100 Bioanalyzer. Ribosomal RNAs should appear as sharp bands or peaks. The apparent ratio of 28S rRNA to 18S rRNA should be approximately 2:1. If the ribosomal bands or peaks of a specific sample are not sharp, but appear as a smear towards smaller sized RNAs, it is likely that the sample suffered major degradation either before or during RNA purification. As a useful measure of RNA integrity, the QIAxcel® Advanced system and the Agilent 2100 Bioanalyzer provide an RNA integrity score (RIS) and an RNA integrity number (RIN), respectively. Ideally, the value should be close to 10, but in many cases (particularly with tissue samples), RNA quality is greatly influenced by how well the original sample was preserved.

* When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, consult the appropriate safety data sheets (SDSs), available from the product supplier.

Appendix C: RNA Cleanup and Concentration

The RNeasy Plus Micro Kit can be used to clean up crude RNA samples. For RNA samples that were purified by phenol/chloroform extraction and will be analyzed using sensitive methods, such as real-time RT-PCR, cleanup of the RNA with the RNeasy Plus Micro Kit, ensures effective removal of contaminants, such as genomic DNA and phenol, which can interfere with transcript quantification. The kit also has the additional benefit of concentrating RNA samples, enabling the use of more RNA template per reaction and/or smaller reaction volumes.

Procedure

1. Add 350 μ l Buffer RLT Plus to a maximum of 50 μ l of crude RNA, or dissolve a crude RNA pellet in 350 μ l Buffer RLT Plus. Mix well by pipetting.
2. Proceed to step 4 of the protocol for animal and human cells on page 21.

Appendix D: Purification of Total RNA Containing Small RNAs from Cells

The following procedure allows the purification of total RNA containing small RNAs, such as miRNA from animal and human cells.

Reagents to be supplied by user

- Ethanol (100%)*

Important points before starting

- Perform all steps of the procedure at room temperature (15–25°C). During the procedure, work quickly.
- Perform all centrifugation steps at 20–25°C in a standard microcentrifuge. Ensure that the centrifuge does not cool below 20°C.

Things to do before starting

- Buffer RPE is supplied as a concentrate. Before using for the first time, add 4 volumes of ethanol (96–100%) as indicated on the bottle to obtain a working solution.

Procedure

Carry out the protocol starting on page 21 up to and including step 4. Instead of performing steps 5–11 (purification of total RNA >200 nucleotides), follow steps 1–6 below (purification of total RNA containing small RNAs).

* When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, consult the appropriate safety data sheets (SDSs), available from the product supplier.

1. Add 1.5 volumes (usually 525 μ l) of 100% ethanol to the flow-through from the gDNA Eliminator spin column, and mix well by pipetting. Do not centrifuge. Proceed immediately to step D2.

Note: The volume of 100% ethanol to add may be less than 525 μ l if some lysate was lost during homogenization and DNA removal.

Note: Precipitates may be visible after addition of ethanol, but this does not affect the procedure.

2. Transfer 700 μ l of the sample, including any precipitate that may have formed, to an RNeasy MinElute spin column placed in a 2 ml collection tube (supplied). Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm). Discard the flow-through.* Repeat step D2 until the whole sample has passed through the membrane. Discard the flow-through each time.

Reuse the collection tube in step D3.

3. Add 500 μ l Buffer RPE to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 15 s at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the flow-through.

Reuse the collection tube in step D4.

Note: Buffer RPE is supplied as a concentrate. Ensure that ethanol is added to Buffer RPE before use (see “Things to do before starting”).

* Flow-through contains Buffer RLT Plus and is therefore not compatible with bleach. See page 6 for safety information.

-
4. Add 500 μ l Buffer RPE to the RNeasy MinElute spin column. Close the lid gently, and centrifuge for 2 min at $\geq 8000 \times g$ ($\geq 10,000$ rpm) to wash the spin column membrane. Discard the collection tube with the flow-through.

Note: After centrifugation, carefully remove the RNeasy MinElute spin column from the collection tube so that the column does not contact the flow-through. Otherwise, carryover of ethanol will occur.

5. Place the RNeasy MinElute spin column in a new 2 ml collection tube (supplied). Open the lid of the spin column, and centrifuge at full speed for 5 min. Discard the collection tube with the flow-through.

To avoid damage to their lids, place the spin columns into the centrifuge with at least one empty position between columns. Orient the lids so that they point in a direction opposite to the rotation of the rotor (e.g., if the rotor rotates clockwise, orient the lids counterclockwise).

It is important to dry the spin column membrane, since residual ethanol may interfere with downstream reactions. Centrifugation with the lids open ensures that no ethanol is carried over during RNA elution.

6. Place the RNeasy MinElute spin column in a new 1.5 ml collection tube (supplied). Add 14 μ l RNase-free water directly to the center of the spin column membrane. Close the lid gently, and centrifuge for 1 min at full speed to elute the RNA (total RNA containing small RNAs).

As little as 10 μ l RNase-free water can be used for elution if a higher RNA concentration is required, but the yield will be reduced by approximately 20%. Do not elute with less than 10 μ l RNase-free water, as the spin column membrane will not be sufficiently hydrated.

The dead volume of the RNeasy MinElute spin column is 2 μ l: elution with 14 μ l RNase-free water results in a 12 μ l eluate.

For real-time RT-PCR with the purified RNA, QIAGEN offers the miScript System, which allows detection of hundreds of miRNAs from a single cDNA synthesis reaction

Appendix E: Acetone Precipitation of Protein from Lysates

The following procedure describes how to recover denatured protein by acetone precipitation from lysates of cells and easy-to-lyse tissues.

Reagents to be supplied by user

- Ice
- Acetone*
- **Optional:** Ethanol*
- Buffer* for downstream application (e.g., loading buffer for SDS-PAGE gel)

Important points before starting

- **Do not use trichloroacetic acid (TCA) to precipitate protein from Buffer RLT Plus lysates.**
This buffer contains guanidine thiocyanate, which can form highly reactive compounds when combined with acidic solutions.

Procedure

Bind total RNA to the RNeasy MinElute spin column as described in the cell protocol (from page 21, steps 1–6) or the tissue protocol (from page 30, steps 1–5). Then follow steps 1–5 below to precipitate protein from the flow-through.

7. Add 4 volumes of ice-cold acetone to the flow-through from the RNeasy MinElute spin column.
8. Incubate for 30 min on ice or at -30°C to -15°C .

* When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, consult the appropriate safety data sheets (SDSs), available from the product supplier.

9. Centrifuge for 10 min at full speed in a benchtop centrifuge. Discard the supernatant and air-dry the pellet.*

10. **Optional:** Wash the pellet with 100 µl ice-cold ethanol and air-dry.

Do not overdry the pellet as this may make resuspension more difficult.

11. Resuspend the pellet in the buffer for your downstream application.

Sodium dodecyl sulfate (SDS) causes guanidine salts to precipitate. In case the pellet contains traces of guanidine thiocyanate, load the sample onto an SDS-PAGE gel immediately after heating for 7 minutes at 95°C.

* Supernatant contains guanidine thiocyanate and is therefore not compatible with bleach. See page 6 for safety information.

Ordering Information

Product	Contents	Cat. no.
RNeasy Plus Micro Kit (50)	50 RNeasy MinElute Spin Columns, 50 gDNA Eliminator Mini Spin Columns, Collection Tubes, Carrier RNA, RNase-Free Reagents and Buffers	74034
RNeasy Plus Mini Kit (50)*	50 RNeasy Mini Spin Columns, 50 gDNA Eliminator Mini Spin Columns, Collection Tubes, RNase-Free Reagents and Buffers	74134
Accessories		
RNAprotect Cell Reagent (250 ml)	250 ml RNAprotect Cell Reagent	76526
RNAprotect Tissue Reagent (50 ml)	50 ml RNAprotect Tissue Reagent for stabilization of RNA in 25 x 200 mg tissue samples	76104
RNAprotect Tissue Reagent (250 ml)	250 ml RNAprotect Tissue Reagent for stabilization of RNA in 125 x 200 mg tissue samples	76106
RNAprotect Tissue Tubes (50 x 1.5 ml)	For stabilization of RNA in 50 x 50 mg tissue samples: 50 screw-top tubes containing 1.5 ml RNAprotect Tissue Reagent each	76154
RNAprotect Tissue Tubes (20 x 5 ml)	For stabilization of RNA in 20 x 500 mg tissue samples: 20 screw-top tubes containing 5 ml RNAprotect Tissue Reagent each	76163
Allprotect Tissue Reagent (100 ml)	For stabilization of DNA/RNA/protein in 50 x 200 mg tissue samples: 100 ml Allprotect Tissue Reagent, Allprotect Reagent Pump	76405
Collection Tubes (2 ml)	1000 x 2 ml Collection Tubes	19201
QIAshredder (50)	50 disposable cell-lysate homogenizers	79654
QIAshredder (250)	250 disposable cell-lysate homogenizers	79656

Product	Contents	Cat. no.
TissueRuptor II	Handheld rotor–stator homogenizer, 5 TissueRuptor Disposable Probes	Varies†
TissueRuptor Disposable Probes (25)	25 nonsterile plastic disposable probes for use with the TissueRuptor II	990890
TissueLyser II	Universal laboratory mixer-mill disruptor	Varies†
TissueLyser Adapter Set 2 x 24	2 sets of Adapter Plates and 2 racks for use with 2 ml microcentrifuge tubes on the TissueLyser II	69982
Stainless Steel Beads, 5 mm (200)	Stainless Steel Beads, suitable for use with the TissueLyser system	69989
TissueLyser Single-Bead Dispenser, 5 mm	For dispensing individual beads (5 mm diameter)	69965
QIAcube Connect — for fully automated nucleic acid extraction with QIAGEN spin-column kits		
QIAcube Connect‡	Instrument, connectivity package, 1-year warranty on parts and labor	Inquire
Starter Pack, QIAcube	Filter-tips, 200 µl (1024), 1000 µl filter-tips (1024), 30 ml reagent bottles (12), rotor adapters (240), elution tubes (240), rotor adapter holder	990395
Related products		
AllPrep® DNA/RNA Micro Kit — for simultaneous purification of DNA and RNA from the same small cell or tissue sample		
AllPrep DNA/RNA Micro Kit (50)	50 AllPrep DNA Mini Spin Columns, 50 RNeasy MinElute Spin Columns, Collection Tubes, RNase-Free Reagents and Buffers	80284
RNeasy FFPE Kit — for purification of high yields of usable RNA from FFPE tissue sections		
RNeasy FFPE Kit (50)	50 RNeasy MinElute Spin Columns, 50 gDNA Eliminator Mini Spin Columns, Collection Tubes, RNase-Free Reagents and Buffers	74404

Product	Contents	Cat. no.
RNeasy MinElute Cleanup Kit — for RNA cleanup and concentration with small elution volumes		
RNeasy MinElute Cleanup Kit (50)	50 RNeasy MinElute Spin Columns, Collection Tubes, RNase-Free Reagents and Buffers	74204
QuantiTect Whole Transcriptome Kit — for unlimited real-time PCR analysis from precious RNA samples		
QuantiTect Whole Transcriptome Kit (25) [§]	For 25 x 50 µl whole-transcriptome-amplification reactions: T-Script Enzyme and Buffer, Ligation Enzymes, Reagent and Buffer, and REPL-g [®] DNA Polymerase and Buffer	207043
Sensiscript RT Kit — for reverse transcription using less than 50 ng RNA per reaction		
Sensiscript RT Kit (50) [§]	For 50 x 20 µl reverse-transcription reactions: Sensiscript Reverse Transcriptase, 10x Buffer RT, dNTP Mix, RNase-Free Water	205211
QuantiNova Reverse Transcription Kit for fast cDNA synthesis and reproducible real-time two-step RT-PCR		
QuantiNova Rev. Transcription Kit (10) [§]	For 10 x 20 µl reactions: 20 µl 8x gDNA Removal Mix, 10 µl Reverse Transcription Enzyme, 40 µl Reverse Transcription Mix (containing RT primers), 20 µl Internal Control RNA, 1.9 ml RNase-Free Water	205410
QuantiNova[®] SYBR[®] Green PCR Kit — or unparalleled results using SYBR Green-based qPCR		
QuantiNova SYBR [®] Green PCR Kit (2500)	For 2500 x 20 µl reactions: 15 x 1.7 ml 2x QuantiNova SYBR Green PCR Master Mix, 5 x 500 µl QuantiNova Yellow Template Dilution Buffer, 5 X 1 ml QN ROX Reference Dye, 5 x 1.9 ml Water	208056
QuantiNova SYBR[®] Green RT-PCR Kit — for one-step qRT-PCR using SYBR[®] Green I for gene expression analysis		

Product	Contents	Cat. no.
QuantiNova® SYBR Green RT-PCR Kit (500)	For 500 x 20 µl reactions: 3 x 1.7 ml QuantiNova SYBR Green RT-PCR Master Mix, 100 µl QuantiNova SYBR Green RT Mix, 100 µl Internal Control RNA, 500 µl Yellow Template Dilution Buffer, 1 ml ROX Reference Dye, 2 x 1.9 µl RNase-Free Water	208154
QuantiNova Probe PCR Kits – for highly sensitive, specific, and ultrafast, probe-based real-time PCR		
QuantiNova Probe PCR Kit (500)	For 500 x 20 µl reactions: 3 x 1.7 ml 2x QuantiNova Probe PCR Master Mix, 1 ml QN ROX Reference Dye, 500 µl QuantiNova Yellow Template Dilution Buffer, 1.9 ml Water	208254
QuantiNova Probe PCR Kits – for highly sensitive, specific, and ultrafast, probe-based real-time PCRs		
QuantiNova Probe RT-PCR Kit (500)	For 500 x 20 µl reactions: 3 x 1.7 ml QuantiNova Probe RT-PCR Master Mix, 100 µl QuantiNova Probe RT Mix, 100 µl Internal Control RNA, 500 µl Yellow Template Dilution Buffer, 1 ml ROX Reference Dye, 2 x 1.9 µl RNase-Free Water	208354
QuantiNova Probe PCR Kit (500)	For 500 x 20 µl reactions: 3 x 1.7 ml 2x QuantiNova Probe PCR Master Mix, 1 ml QN ROX Reference Dye, 500 µl QuantiNova Yellow Template Dilution Buffer, 1.9 ml Water	208354

* For total RNA purification from up to 1×10^7 cells or 30 mg tissue.

† Visit www.qiagen.com/automation to find out more about the TissueRuptor II and TissueLyser II and to order.

‡ All QIAcube Connect instruments are provided with a region-specific connectivity package, including tablet and equipment necessary to connect to the local network. Further, QIAGEN offers comprehensive instrument service products, including service agreements, installation, introductory training and preventive subscription. Contact your local sales representative to learn about your options.

§ Other kit sizes available; please inquire.

¶ For all instruments from Applied Biosystems except the Applied Biosystems 7500.

** For the Applied Biosystems 7500 and instruments from Bio-Rad/MJ Research, Cepheid, Corbett Research, Eppendorf, Roche and Stratagene.

For up-to-date licensing information and product-specific disclaimers, see the respective QIAGEN kit handbook or user manual. QIAGEN kit handbooks and user manuals are available at www.qiagen.com or can be requested from QIAGEN Technical Services or your local distributor.

Document Revision History

Date	Changes
January 2020	Updated text, ordering information and intended use for QIAcube Connect.
November 2020	Updated branding of RNA protection products.

Trademarks: QIAGEN[®], Sample to Insight[®], QIAcube[®], QIAxcel[®], AllPrep[®], Allprotect[®], GeneGlobe[®], MinElute[®], Quantifast[®], QuantiNova[®], QuantiTect[®], REPLiG[®], RNAprotect[®], RNeasy[®], Rotor-Gene[®], Sensiscript[®] (QIAGEN Group); Agilent[®] (Agilent Technologies, Inc.); Applied Biosystems[®] (Applied Biosystems Corporation or its subsidiaries); FACS[®] (Becton, Dickinson and Company); Leica[®] (Leica Microsystems GmbH); SYBR[®] (Molecular Probes, Inc.). Registered names, trademarks, etc. used in this document, even when not specifically marked as such, are not to be considered unprotected by law.

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