

MACHEREY-NAGEL

# User manual



## RNA isolation






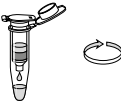
- NucleoSpin® RNA Plus XS

July 2025 2022 / Rev. 8

# RNA isolation

## Protocol at a glance (Rev. 08)

NucleoSpin® RNA Plus XS

<b>1 Homogenize and lyse sample</b>		100 µL LB1 Homogenize 100 µL LB2
<b>2 Remove gDNA and filtrate lysate</b>		100 x g, 2 min 11,000 x g, 10 s
<b>3 Adjust binding conditions</b>		150 µL BSXS Mix
<b>4 Bind RNA</b>		Load lysate 300 x g, 1 min 11,000 x g, 10 s
<b>5 Wash and dry silica membrane</b>		1 <sup>st</sup> wash 100 µL MDB 11,000 x g, 10 s 2 <sup>nd</sup> wash 500 µL WB2 11,000 x g, 10 s 3 <sup>rd</sup> wash 200 µL WB2 < 20,000 x g, 2 min
<b>6 Elute RNA</b>		10–20 µL RNase-free H <sub>2</sub> O 11,000 x g, 1 min

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# 1 Components

## 1.1 Kit contents

NucleoSpin® RNA Plus XS			
REF	10 preps 740990.10	50 preps 740990.50	250 preps 740990.250
Lysis Buffer LB1	6 mL	6 mL	30 mL
Lysis Buffer LB2	6 mL	6 mL	30 mL
Binding Solution BSXS	10 mL	10 mL	50 mL
Wash Buffer MDB	10 mL	10 mL	30 mL
Wash Buffer WB2 (concentrate)	6 mL	12 mL	50 mL
RNase-free H <sub>2</sub> O	13 mL	13 mL	13 mL
NucleoSpin® gDNA Removal Column XS (yellow ring)	10	50	250
NucleoSpin® RNA Plus XS Column (light blue ring plus collection tube)	10	50	250
Collection Tube (2 mL)	20	100	500
Collection Tube (1.5 mL)	10	50	250
User manual	1	1	1

## 1.2 Reagents, consumables, and equipment to be supplied by user

### Reagents

- 96–100 % ethanol (to prepare Wash Buffer WB2, non-denatured ethanol recommended)

### Consumables

- 1.5 mL or 2.0 mL microcentrifuge tubes or PCR reaction tubes (to prepare sample lysate)
- Disposable micro pipette (optional, for sample homogenisation)
- Sterile RNase-free tips
- NucleoSpin® Bead Tubes Type A or C (optional, please see 6.3 Ordering information)

### Equipment

- Manual pipettes
- Vortex mixer
- Centrifuge for microcentrifuge tubes
- Equipment for sample disruption and homogenization (see section 2.3)
- Personal protection equipment (e.g., lab coat, gloves, goggles)

*Note: Additional reducing agents (e.g.,  $\beta$ -mercaptoethanol, DTT, TCEP) are not required for NucleoSpin® RNA Plus XS preparations.*

## 1.3 RNase-free work environment

Kit components have been tested to ensure they are RNase-free. However, a RNase-free working environment is also a critical factor for performing successful RNA isolation and handling. Therefore, general recommendations to avoid RNase contamination should be followed:

- Maintain a separate area, RNase-free pipettes, and materials when working with RNA.
- Wear gloves when handling RNA and reagents to avoid contact with skin, which is a source of RNases. Change gloves frequently.
- Use sterile RNase-free plastic tubes. Collection Tubes (2 mL, for column flowthrough and 1.5 mL for elution) are provided in the kit. Tubes for lysate preparation have to be supplied by user.
- Use RNase-free water supplied with the kit for elution.
- Keep all kit components sealed when not in use and all tubes tightly closed when possible.

## 1.4 About this user manual

It is strongly recommended reading the detailed protocol sections of this user manual if the **NucleoSpin® RNA Plus XS** kit is used for the first time. Experienced users, however, may refer to the Protocol at a glance instead. The Protocol at a glance is designed to be used only as a supplemental tool for quick referencing while performing the purification procedure.

All technical literature is available on the internet at **[www.mn-net.com](http://www.mn-net.com)**.

Please contact Technical Service regarding information about changes of the current user manual compared to previous revisions.

## 2 Product description

### 2.1 The basic principle

The **NucleoSpin® RNA Plus XS** kit is designed to purify RNA from small amounts of a variety of cell and tissue types. This kit supplies the NucleoSpin® gDNA Removal Column XS, a spin column that quickly and effectively removes genomic DNA without the need of DNase digestion.

One of the most important aspects during the isolation of RNA is to prevent degradation of the RNA. Cells and tissues are first lysed by incubation in a chaotropic ion Lysis Buffer (LB1 + LB2), which immediately inactivates RNases. The lysate is added to the NucleoSpin® gDNA Removal Column XS (yellow rings) to clarify the lysate and to remove contaminating gDNA. After the addition of the Binding Solution BSXS to the flowthrough, the RNA is bound to the **NucleoSpin® RNA Plus XS** Column (light blue rings). Subsequent wash steps remove salts, metabolites, and macromolecular cellular components. High quality RNA is eluted with RNase-free H<sub>2</sub>O.

The RNA preparation using **NucleoSpin® RNA Plus XS** kits can be performed at room temperature.

The eluate should be treated with care because RNA is very sensitive to trace contaminations of RNases, often found on general lab ware, fingerprints, and dust. Keep RNA frozen at -20 °C for short term or -70 °C for long term storage to ensure RNA stability.

### 2.2 Kit specifications

- **NucleoSpin® RNA Plus XS** kits are recommended for the isolation of RNA from small amounts of cultured cells and tissue. The **NucleoSpin® RNA Plus XS** kits allows purification of high quality RNA. The RNA  $A_{260}/A_{280}$  ratio generally exceeds 1.9 (measured in TE buffer, pH 7.5).
- The isolated RNA is ready to use in diverse downstream applications.
- RNA isolated with the **NucleoSpin® RNA Plus XS** kit is of high integrity. RIN (RNA Integrity Number) or RQN (RNA Quality Number) of RNA isolated from fresh high quality sample material (e.g., eukaryotic cells, or fresh mouse liver) generally exceeds 8. However, RNA integrity strongly depends on the sample quality and a sufficient RNA concentration.
- RNA molecules isolated with **NucleoSpin® RNA Plus XS** are longer than approximately 100 nucleotides. Thus, the **NucleoSpin® RNA Plus XS** kit provides an enrichment of mRNA and rRNA. RNA isolated with the **NucleoSpin® RNA Plus XS** kits may contain minute amounts of genomic DNA due to carry-over from the NucleoSpin® gDNA Removal Column XS. The probability of DNA detection with PCR increases with:
  1. The number of DNA copies per preparation: single copy target < plasmid / mitochondrial target < plasmid transfected into cells.
  2. decreasing PCR amplicon size.

**Table 1: Kit specifications at a glance**

Parameter	NucleoSpin® RNA Plus XS
Technology	Two column silica membrane system: 1. XS column for DNA removal 2. XS column for RNA isolation
Use	For research use only
Format	Mini spin column – XS design
Processing	Manual handling and centrifugation
Sample material	< 10 <sup>5</sup> cultured cells < 5 mg tissue
Fragment size	> 100 nt
Typical yield	10 <sup>5</sup> HeLa cells: ca. 500–2000 ng 10 <sup>4</sup> HeLa cells: ca. 50–200 ng 10 <sup>3</sup> HeLa cells: ca. 5–20 ng 10 <sup>2</sup> HeLa cells: ca. 0.5–2 ng 10 <sup>1</sup> HeLa cells: ca. 0.05–0.2 ng 1 HeLa cell: ca. 0.005–0.02 ng  5000 µg liver: ca. 300–500 ng 500 µg liver: ca. 300–500 ng 50 µg liver: ca. 100–250 ng 5 µg liver: ca. 25–70 ng 0.5 µg liver: ca. 2,5–8 ng 0.05 µg liver: ca. 0,25–1,2 ng 0.005 µg liver: 0,025–0,2 ng
A <sub>260</sub> /A <sub>280</sub>	1.9–2.2*
A <sub>260</sub> /A <sub>230</sub>	1.5–2.5*
Typical RIN (RNA Integrity Number)	> 8*
Elution volume	5–30 µL
Preparation time	18 min/6 preps
Binding capacity	110 µg

\*Quality ratio determination strongly depends on a sufficient amount of RNA measured. Make sure to use sufficient amount of RNA that has been validated to enable meaningful ratio determination!

## 2.3 Handling, preparation, and storage of starting materials

The maximum amount of sample material that can be used with the **NucleoSpin® RNA Plus XS** kit depends on type of sample and its RNA and DNA content.

Maximal amount of sample material to be used per preparation (approximate values):

- Eukaryotic cells (e.g., HeLa cells):  $10^5$  cells
- Animal tissue: 5 mg (wet weight) e.g. liver, kidney, or 1 mg spleen (wet weight)

### Sample storage and RNase inhibition

RNases rapidly degrade RNA within the samples if samples are not protected from RNase activity after harvest.

- Use freshly harvested sample for immediate lysis and RNA purification.
- Store samples in lysis buffer after disruption at  $-70\text{ °C}$  for up to one year, at  $4\text{ °C}$  for up to 24 hours or at room temperature for up to several hours. Samples frozen in lysis buffer should be thawed as followed before starting with the isolation of RNA. Place the frozen tubes in a water bath with lukewarm water (maximum  $37\text{ °C}$ ) or in a thermocycler set to  $37\text{ °C}$  (shaking). Stay with the tubes and check them every 30 seconds until fully thawed. As soon as the samples are thawed, promptly remove the tubes from the bath and place them at room temperature. You can then proceed with step 2 of the standard protocol.
- Flash freeze sample in liquid  $\text{N}_2$  immediately upon harvest and store at  $-70\text{ °C}$ . Frozen samples are stable up to 6 months. Mortar and pestle can be used to pulverize the sample in a frozen state. Make sure that the sample does not thaw prior to contact with lysis buffer.
- Submerge and store samples in NucleoProtect® RNA or RNAlater® stabilization solution. Make sure to allow complete permeation of the sample with the stabilization solution before freezing it. Remove excess stabilization solution from the sample prior to RNA isolation according to the stabilization solution user manual.

### Disruption and homogenisation of sample material

#### Adherent cells

Completely aspirate cell culture medium and immediately add Lysis Buffer LB1 to the cell culture dish. Avoid incomplete removal of the cell culture medium in order to allow full lysis activity of the Lysis Buffer. For lysis vortex cells vigorously.

#### To trypsinize adherent growing cells:

Aspirate cell culture medium, and add an equal amount of PBS in order to wash the cells. Aspirate PBS. Add 0.1–0.3% trypsin in PBS and incubate for an appropriate time to detach the cells from the dish surface. After cell detachment, add cell culture medium, transfer cells to an appropriate tube (not supplied), and pellet by centrifugation for 5 min at  $300 \times g$ . Remove supernatant and continue with the addition of lysis buffer to the cell pellet.

#### Cells in suspension

Spin down cells for 5 min at  $300 \times g$ , remove the supernatant completely and directly add Lysis Buffer LB1 to the cell pellet. Vortex vigorously for resuspension and lysis.

## Tissue samples

- Disruption can be performed by NucleoSpin® Bead Tubes Type A or C (see ordering information). In combination with a Vortex-Genie® 2 the tubes can be mounted on a MN Bead Tube Holder (see ordering information). Add sample and Lysis Buffer LB1 to the bead tube and agitate for 5 min. Alternatively, the NucleoSpin® Bead Tubes A can be used in a bead mill. NucleoSpin® Bead Tubes Type C are not recommended to use with a bead mill. This might be necessary for some tissue types (e.g., lung, heart tissue). Optimal time and frequency have to be determined experimentally. It is recommended to reduce the amount of beads in the tube from approximately 400 µL to approximately 70 µL–100 µL beads in order to enable efficient lysate withdrawal from the Bead Tube.
- Microtube pestles can be used to homogenize frozen material or fresh material in Lysis Buffer LB1 directly in a microcentrifuge tube.

For tissue samples the lysate should be clarified by centrifugation for 3 min at 5000 x g. The supernatant is ready to use for **NucleoSpin® RNA Plus XS** protocol. For cultivated cells a clarification is typically not necessary.

## 2.4 Elution procedures

Elution volumes in the range of 5–30 µL are recommended. The default elution volume is 10–20 µL.

## 2.5 Stability of isolated RNA

Eluted RNA should always be kept on ice during work for optimal stability. Contamination with almost omnipresent RNases (general lab ware, fingerprints, dust) may lead to degradation of isolated RNA. For short term storage, freeze RNA at –20 °C, for long term storage freeze at –70 °C.

### 3 Storage conditions and preparation of working solutions

**Attention:**

*Buffers LB1, LB2, and MDB contain chaotropic salt. Wear gloves and goggles!*

*CAUTION: Buffers LB1, LB2, and MDB contain chaotropic salt which can form highly reactive compounds when combined with bleach (sodium hypochlorite). DO NOT add bleach or acidic solutions directly to the sample-preparation waste.*

- All kit components should be stored at 15 – 25 °C and are stable until: see package label. Storage at lower temperatures may cause precipitation of salts.

Before starting any **NucleoSpin® RNA Plus XS** protocol, prepare the following:

- **Wash Buffer WB2:** Add the indicated volume of 96 – 100 % ethanol (see table below) to Buffer WB2 Concentrate. Mark the label of the bottle to indicate that ethanol was added. Wash Buffer WB2 can be stored at 15 – 25 °C for at least one year.
- If desired, aliquot RNase-free water into RNase-free 1.5 mL tubes (not provided) in order to avoid accidental RNase contamination of the water by repeated opening of the bottle.

NucleoSpin® RNA Plus XS			
REF	10 preps 740990.10	50 preps 740990.50	250 preps 740990.250
Wash Buffer WB2 (concentrate)	6 mL Add 24 mL ethanol	12 mL Add 48 mL ethanol	50 mL Add 200 mL ethanol

## 4 Safety instructions

When working with the **NucleoSpin® RNA Plus XS** kit wear suitable protective clothing (e.g., lab coat, disposable gloves, and protective goggles). For more information consult the appropriate Material Safety Data Sheets (MSDS available online at [www.mn-net.com/msds](http://www.mn-net.com/msds)).



Caution: Guanidinium thiocyanate in buffer LB1 and buffer LB2 can form highly reactive compounds when combined with bleach! Thus, do not add bleach or acidic solutions directly to the sample preparation waste.

The waste generated with the **NucleoSpin® RNA Plus XS** kit has not been tested for residual infectious material. A contamination of the liquid waste with residual infectious material is highly unlikely due to strong denaturing lysis buffer treatment but it cannot be excluded completely. Therefore, liquid waste must be considered infectious and should be handled and discarded according local safety regulations.

### 4.1 Disposal

Dispose hazardous, infectious or biologically contaminated materials in a safe and acceptable manner and in accordance with all local and regulatory requirements.

## 5 NucleoSpin® RNA Plus XS Protocol

### Before starting the preparation:

- Check if Wash Buffer WB2 was prepared according to section 3.

### 1 Homogenize and lyse sample

Disrupt and clarify the sample according to one of the methods described in section 2.3 by using **100 µL Lysis Buffer LB1**.



100 µL LB1  
Homogenize  
100 µL LB2

#### Cells

Add LB1 and vortex vigorously.

#### Tissue

Homogenize in presence of LB1 by using microtube pestle or bead beating.

Clarify tissue lysates by centrifugation for **3 min** at **5000 x g**. If a visible pellet is observed, it is recommended to transfer the supernatant into a fresh tube and discard the pellet.

*Note:* Lysis tube is not included in the kit. Addition of reducing agent (e.g.,  $\beta$ -mercaptoethanol, DTT, or TCEP) is not necessary.

Add **one volume Lysis Buffer LB2** to the lysate (typically 100 µL) and mix.

*Note:* 100 µL LB1 is sufficient for lysis of samples < 20 µL in volume (e.g., 1 mg tissue or low titer cell suspensions). For processing of larger samples (e.g., cell suspensions of > 20 µL) or if a larger lysate volume is desired (e.g., for bead beating), increase the volumes of Lysis Buffer LB1, LB2, and Binding solution BSXS proportionally. Be aware that 180 µL LB1 (+ 180 µL LB2 + 270 µL BSXS) should not be exceeded due to the restricted volume of the column and collection tube. If higher volumes of LB1 (REF 740368.30) and LB2 (REF 740369.30) are used, additional buffer needs to be ordered separately, see 6.3 Ordering information.

**2 Remove gDNA and filtrate lysate**

Place a **NucleoSpin® gDNA Removal Column XS** (yellow ring) in a Collection Tube (2 mL, provided), **transfer** the homogenized **lysate** (~200 µL) to the column, and centrifuge for **2 min** at **100 x g** followed by **10 s** at **11,000 x g**.



100 x g,  
2 min



11,000 x g,  
10 s

Discard the column and **continue with the flowthrough**.

*Note: Ensure that no liquid remains on the column membrane after centrifugation. If necessary, repeat the centrifugation until all liquid has passed through the membrane. The low g-centrifugation allows efficient DNA removal.*

**3 Adjust RNA binding conditions**

Add **150 µL Binding Solution BSXS** (1.5 volumes relative to LB1) to the flowthrough and mix well by moderate vortexing or by pipetting up and down several times.



150 µL BSXS  
Mix

*Note: If mixing is done by vortexing, be careful in order to avoid spilling, because the Collection Tube does not contain a lid.*

**4 Bind RNA**

**Transfer** the whole **lysate** (~350 µL) to the **NucleoSpin® RNA Plus XS Column** (light blue ring) preassembled with a Collection Tube.



load lysate  
300 x g,  
1 min

Centrifuge for **1 min** at **300 x g** followed by **10 s** at **11,000 x g**.



11,000 x g,  
10 s

*Note: The flowthrough may remain in the collection tube.*

**5 Wash and dry silica membrane****1<sup>st</sup> wash**

Add **100 µL Wash Buffer MDB** onto the column.

Centrifuge for **10 s** at **11,000 x g**.

Discard the flowthrough with collection tube and place the column into a new 2 mL Collection Tube (provided).



100 µL MDB

11,000 x g,  
10 s**2<sup>nd</sup> wash**

Add **500 µL Buffer WB2** onto the column.

Centrifuge for **10 s** at **11,000 x g**.

*Discard the flowthrough and reuse the collection tube.*



500 µL WB2

11,000 x g,  
10 s**3<sup>rd</sup> wash**

Add **200 µL Buffer WB2** onto the column.

Centrifuge for **2 min** at **full speed** (< 20,000 x g) to dry the membrane.

Place the column into a **Collection Tube** (1.5 mL, provided).

*Note: If for any reason, the liquid level in the Collection Tube has reached the column after centrifugation, discard flowthrough and centrifuge again.*



200 µL WB2

< 20,000 x g,  
2 min**6 Elute RNA**

Add **10–20 µL RNase-free H<sub>2</sub>O** and centrifuge **1 min** at **11,000 x g**.

*Note: Elution volume may be varied from 5 µL to 30 µL.*

10–20 µL  
RNase-free  
H<sub>2</sub>O11,000 x g,  
1 min

## 6 Appendix

### 6.1 Removal of DNA

In case samples with high initial DNA content are analyzed by downstream applications highly sensitive towards DNA contamination, an additional DNA digest might be required. The protocol for DNase treatments is given below.

#### Protocol A: DNA digestion in solution

##### 1 Digest DNA (Reaction setup)

Add **1 µL Reaction Buffer for rDNase** and **0.1 µL rDNase** to **10 µL** eluted RNA.

*Note: Alternatively, premix 100 µL Reaction Buffer for rDNase and 10 µL rDNase and add 1/10 volume to one volume of RNA eluate.*

Gently swirl the tube in order to mix the solution. Spin down gently (approx. 1 s at 1,000 x g) to collect every droplet of the solution at the bottom of the tube.

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##### 2 Incubate sample

Incubate for **10 min** at **37 °C**.

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##### 3 Repurify RNA

Repurify RNA with a suitable RNA clean up procedure, for example by use of the NucleoSpin® RNA Clean-up or NucleoSpin® RNA Clean-up XS kits (please see 6.3 Ordering information), or by ethanol precipitation.

##### **Ethanol precipitation, exemplary:**

Add **0.1 volume** of **3 M sodium acetate, pH 5.2** and **2.5 volumes** of **96–100 % ethanol** to **one volume** of sample. Mix thoroughly.

Incubate **several minutes** to **several hours** at **–20 °C** or **4 °C**.

*Note: Choose long incubation times if the sample contains low RNA concentration. Short incubation times are sufficient if the sample contains high RNA concentration.*

Centrifuge for **10 min** at **maximum speed**.

Wash RNA pellet with 70 % ethanol.

Dry RNA pellet and resuspend RNA in RNase-free H<sub>2</sub>O.

## **Protocol B: On column DNA digestion**

### **1 Reconstitution of rDNase**

Add 4 mL Reaction Buffer for rDNase into a rDNase Vial Size F and dissolve the DNase.

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### **2 On-Column digestion**

Follow the purification procedure according to section 5 until the column has been washed with 100  $\mu$ L Wash Buffer MDB (step 5).

Apply **95  $\mu$ L rDNase reaction mixture** directly onto the center of the silica membrane of the column.

Incubate at room temperature for 15 min.

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Continue the procedure 5.1, step 5, by adding 200  $\mu$ L Buffer WB2 onto the column.

## 6.2 Troubleshooting

Problem	Possible cause and suggestions
RNA is degraded/no RNA obtained	<i>RNase contamination</i> <ul style="list-style-type: none"><li>• Create an RNase free working environment. Wear gloves during all steps of the procedure. Change gloves frequently. Use of sterile, disposable polypropylene tubes is recommended. Keep tubes closed whenever possible during the preparation. Glassware should be oven-baked for at least 2 hours at 250 °C before use.</li></ul>
	<i>Insufficient sample quality</i> <ul style="list-style-type: none"><li>• Control sample harvest, storage and lysis. Make sure that samples are harvested, stored, and lysed adequately in order to preserve RNA integrity. See section 2.3 for recommended procedures.</li></ul>
Poor RNA quality or yield	<i>Reagents not applied or restored properly</i> <ul style="list-style-type: none"><li>• Reagents not properly restored. Add the indicated volume of 96 % ethanol to Buffer WB2 Concentrate and mix.</li><li>• Sample and reagents have not been mixed completely. Always vortex vigorously after each reagent has been added.</li><li>• No Binding Solution BSXS has been added after lysis. Binding of RNA to the silica membrane is only effective in the presence of Binding Solution.</li></ul>
	<i>Kit storage</i> <ul style="list-style-type: none"><li>• Store kit components at room temperature. Storage at low temperatures may cause salt precipitation.</li><li>• Keep bottles tightly closed in order to prevent evaporation or contamination.</li></ul>

Fine-tune your nucleic acid yield and purity



[www.mn-net.com/nucleic-acid-fine-tune](http://www.mn-net.com/nucleic-acid-fine-tune)

*Ionic strength and pH influence  $A_{260}$  absorption as well as ratio  $A_{260}/A_{280}$*

For adsorption measurement, use 5 mM Tris pH 8.5 as diluent. Please see also:

- Manchester, K L. 1995. Value of  $A_{260}/A_{280}$  ratios for measurement of purity of nucleic acids. *Biotechniques* 19, 208–209.
- Wilfinger, W W, Mackey, K and Chomczynski, P. 1997. Effect of pH and ionic strength on the spectrophotometric assessment of nucleic acid purity. *Biotechniques* 22, 474–481.

Poor RNA quality or yield (continued)

*Sample material*

- Sample material not stored properly. Whenever possible, use fresh material. If this is not possible, flash freeze the samples in liquid  $N_2$ . Samples should always be kept at  $-70\text{ }^\circ\text{C}$ . Never allow tissues to thaw before addition of Lysis Buffer LB1. Perform disruption of samples in liquid  $N_2$ . Alternatively, RNA stabilization solutions may be used to protect RNA from degradation (e.g., NucleoProtect® RNA; see ordering information).
- Insufficient disruption and/or homogenization of starting material. Ensure thorough sample disruption.

*Carry-over of guanidinium thiocyanate*

Low  $A_{260}/A_{230}$  ratio

- Carefully load the lysate to the NucleoSpin® RNA Plus XS Column and try to avoid a contamination of the upper part of the column and the column lid.
- Make sure that a sufficient amount / concentration of RNA is used for quantification so that the  $A_{230}$  value is significantly higher than the background level.
- Measurement of low amount / concentration of RNA will cause unstable  $A_{260}/A_{230}$  ratio values.

*Sample material*

Clogged NucleoSpin® Column / Poor RNA quality or yield

- Too much starting material used. Overloading may lead to decreased overall yield. Reduce amount of sample material or use larger volume of lysis buffer.
- Insufficient disruption and/or homogenization of starting material. Ensure thorough sample disruption and use NucleoSpin® gDNA Removal Column XS for DNA removal and for easy homogenization of disrupted starting material.
- Increase  $g$ -force and centrifugation time if necessary.

*Too much cell material used*

- Reduce quantity of cells or tissue used.

*DNA detection system too sensitive*

Contamination  
of RNA with  
genomic DNA

- The amount of DNA contamination is effectively reduced by the NucleoSpin® gDNA Removal Column XS. However, it can not be guaranteed that the purified RNA is 100 % free of DNA, therefore in very sensitive applications it might still be possible to detect DNA. The probability of DNA detection with PCR increases with:
  - the number of DNA copies per preparation: single copy target < plasmidial / mitochondrial target < plasmid transfected into cells
  - decreasing of PCR amplicon size.
- Use larger PCR targets (e.g., > 500 bp) or intron spanning primers if possible.
- **Use support protocol 6.1 for subsequent rDNase digestion in solution.**

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*Carry-over of ethanol or salt*

Suboptimal  
performance  
of RNA in  
downstream  
experiments

- Do not let the flowthrough touch the column outlet after the second Buffer WB2 wash. Be sure to centrifuge at the corresponding speed for the respective time in order to remove ethanolic Buffer WB2 completely.
- Check if Buffer WB2 has been equilibrated to room temperature before use. Washing at lower temperatures lowers efficiency of salt removal by Buffer WB2.

*Store isolated RNA properly*

- Eluted RNA should always be kept on ice for optimal stability since trace contaminations of omnipresent RNases (general lab ware, fingerprints, dust) will degrade the isolated RNA. For short term storage freeze at -20 °C, for long term storage freeze at -70 °C.

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*Too much sample material*

Too much  
gDNA  
contamination

- A: Use 150 µL LB1 and 50 µL LB2 instead of a 1:1 ratio. This will reduce gDNA further, but RNA yield will decrease. However, the RNA to DNA ratio will improve.
- B: Perform an DNA digestion in solution or on column according to section 6.1

*Store isolated RNA properly*

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*LB2 not applied*

No RNA yield

- For optimal results, lysis should be performed in LB1 and then the lysate should be mixed with on volume LB2. If only LB1 is used, DNA and RNA will bind to the NucleoSpin® gDNA Removal Column XS

### 6.3 Ordering information

Product	REF	Pack of
NucleoSpin® RNA Plus XS	740990.10 / .50 / .250	10 / 50 / 250
NucleoSpin® RNA Plus	740984.10 / .50 / .250	10 / 50 / 250
NucleoZOL	740404.200	200 mL
NucleoSpin® RNA Set for NucleoZOL	740406.50	50
NucleoSpin® RNA Clean-up	740948.10 / .50 / .250	10 / 50 / 250
NucleoSpin® RNA Clean-up XS	740903.10 / .50 / .250	10 / 50 / 250
rDNase Set	740963	1
Collection Tubes	740600	1000
Lysis Buffer LB1	740368.30	30 mL
Lysis Buffer LB2	740369.30	30 mL
NucleoSpin® Bead Tube Type A	740786.50	50
NucleoSpin® Bead Tube Type C	740813.50	50
MN Bead Tube Holder	740469	1
NucleoProtect® RNA	740400.50 / .250 / .500	50 / 250 / 500 mL

## 6.4 Product use restriction / warranty

All MACHEREY-NAGEL products are designed for their intended use only. They are not intended to be used for any other purpose. The description of the intended use of the products can be found in the original MACHEREY-NAGEL product leaflets. Before using our products, please observe the instructions for use and the safety instructions from the respective Material Safety Data Sheet of the product.

This MACHEREY-NAGEL product is carrying documentation stating specifications and other technical information. MACHEREY-NAGEL warrants to meet the stated specifications. The provided warranty is limited to the data specifications and descriptions as given in the original MACHEREY-NAGEL literature. No other statements or representations, written or oral, by MACHEREY-NAGEL's employees, agents or representatives, except written statements signed by a duly authorized officer of MACHEREY-NAGEL are authorized. They should not be relied upon by the customer and are not a part of a contract of sale or of this warranty.

Liability for all possible damages that occur in any connection with our products is limited to the utmost minimum as stated in the general business terms and conditions of MACHEREY-NAGEL in their latest edition which can be taken from the company's website. MACHEREY-NAGEL does not assume any further warranty.

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Tel.: +49 24 21 969-333  
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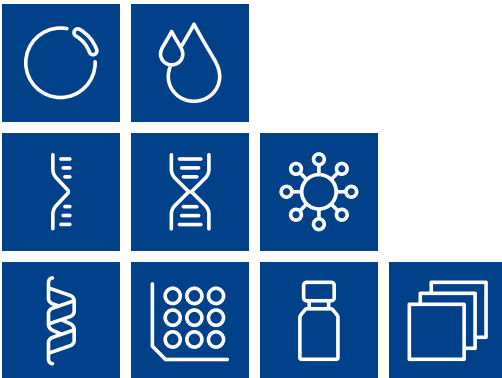
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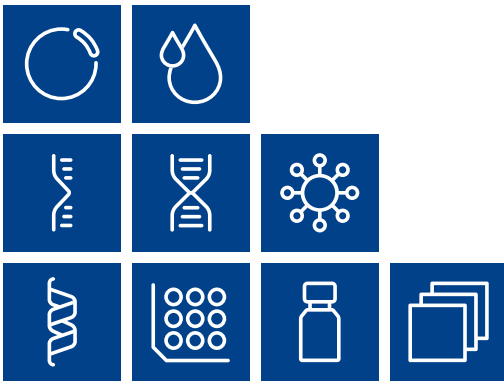
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