NucleoSpin[®] Extract II Kits User Manual

Cat. Nos.636971 636972 636973 PT3814-1 (PR48598) Published 10 September 2004

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I. Kit Contents

TABLE I. NUCLEOSPIN® EXTRACT II			
Cat. No.	10 preps 636971	50 preps 636972	250 preps 636973
Buffer NT	6 ml	30 ml	2 x 75 ml
Buffer NT3 (concentrate)	7 ml	2 x 7 ml	80 ml
Buffer NE	5 ml	15 ml	50 ml
NucleoSpin [®] Extract II columns	10	50	250
NucleoSpin [®] collecting tubes (2ml)	10	50	250
Protocol	1	1	1

II. Product Description

A. The Basic Principle

With the **NucleoSpin® Extract II** method, DNA binds in the presence of chaotropic salt to a silica membrane. The binding mixture is loaded directly onto **NucleoSpin® Extract II** columns. Contaminations like salts and soluble macromolecular components are removed by a simple washing step with ethanolic buffer NT3. Pure DNA is finally eluted under low ionic strength conditions with slightly alkaline buffer NE (5 mM Tris-Cl, pH 8.5).

B. About This User Manual

Experienced users who are performing the purification of PCR* products or DNA fragments from agarose gels using a **NucleoSpin® Extract II** isolation kit may refer to the Protocol-at-a-glance instead of this user manual. The Protocol-at-a-glance is designed to be used only as a supplemental tool for quick referencing while performing the purification procedure. First-time users are strongly advised to read this user manual.

C. Kit Specifications

The **NucleoSpin® Extract II** kit is designed for the purification of DNA from TAE/TBE agarose gels and for direct purification of PCR* products (two applications in one kit).

The **NucleoSpin® Extract II** buffer formulation ensures complete removal of primers from PCR* reactions while small DNA fragments are still bound and purified with high yields

With **NucleoSpin® Extract II** even DNA fragments from PCR reaction buffers rich in various detergents can be purified with high recovery.

The adsorption of DNA to the **NucleoSpin® Extract II** membrane is pH dependent. Optimal yields are achieved by using TAE standard gels or reaction mixtures with pH 6-8.

Standard as well as low melting agarose gels can be used.

The prepared DNA fragments can be used directly in applications like automated fluorescent DNA sequencing, PCR*, or any kind of enzymatic manipulation.

^{*}PCR is patented by Roche Diagnostics

II. Product Description *continued*

TABLE II. KIT SPECIFICATIONS AT A GLANCE			
Parameters	NucleoSpin [®] Extract II		
DNA fragments from agarose gels	Х		
Desalination, removal of enzymes, nucleotides, and/ or labeling reagents like biotin or radioactive ATP etc.	х		
Direct purification of amplified DNA	Х		
Elution volume	15-50 <i>μl</i>		
Binding capacity	15 <i>μg</i>		
Time/prep	10 min for 6 preps		

D. Elution Procedures

- For elution of DNA one of the following solutions can be used: Buffer NE (supplied) / TE buffer, pH 8.5 / distilled water, pH 8.5.
- Note: EDTA in TE buffer may cause problems in subsequent reactions, and the pH of distilled water should be checked before use to avoid lower recovery yields. See Table 1 for correlation between dispensed elution buffer volumes and typical recovery rates for purification of 1-5 µg of PCR* fragments (for gel extraction, recovery rates are approx. 10 % lower).
- With an elution volume of 15 μ / of buffer NE typical recovery rates of 70-95 % are usually obtained for DNA fragments between 50-10,000 bp resulting in highly concentrated eluates (see Table 1). If larger amounts (5-15 μ g) of DNA have to be purified (e.g. from PCR* reactions > 100 μ / or gel slices > 200 mg) elution with at least 50 of buffer NE is recommended. Primers are not bound.
- Yields of larger fragments (> 5-10 kb) can be increased by using prewarmed elution buffer (70°C): For elution, add prewarmed elution buffer NE and incubate for 1-2 min before collecting eluate by centrifugation.For fragments > 10 kb the use of our NucleoTraP®CR kit is recommended.

^{*}PCR is patented by Roche Diagnostics

II. Product Description *continued*

TABLE III. RECOVERY RATES NUCLEOSPIN® EXTRACT II			
Fragment Length	Elution Volume	NucleoSpin [®] Extract II	
65 bp	15 μl 25 μl 50 μl 100 μl	85 % 90 % 95 % 95 %	
400 bp	15 μl 25 μl 50 μl 100 μl	85 % 95 % 100 % 100 %	
700 bp	15 μl 25 μl 50 μl 100 μl	85 % 90 % 95 % 95 %	

II. Product Description *continued*



Figure 1: Use of different elution volumes

A PCR sample with a fragment size of 782 bp were purified according to the standard protocol of **NucleoSpin® Extract II** using different elution volumes as shown. All elution volumes were adjusted to 25 μ l plus 4.5 μ l loading dye. For analysis the mixture was loaded onto a 1% TAE gel. The recovery rates were estimated by use of a fragment ladder.

III. Storage Conditions and Preparation of Working Solutions

Attention:

Buffer NT contains chaotropic salt. Wear gloves and goggles!

• The NucleoSpin® Extract II kit should be stored at room temperature and is stable for up to one year.

Before starting any NucleoSpin® Extract II protocol prepare the following:

Add the indicated volume of 96-100% ethanol to buffer NT3 concentrate.

TABLE IV NUCLEOSPIN® EXTRACT II				
Cat. No.	10 preps	50 preps	250 preps	
	636971	636972	636973	
Buffer NT3	7 ml	2 x 7 ml	80 ml	
	add 28 ml ethanol	add 28 ml ethanol	add 320 ml ethanol	

IV. Safety Instructions – Risk and Safety Phrases

The following components of the NucleoSpin® Extract II kits contain hazardous contents.

Wear gloves and goggles and follow the safety instructions given in this section.

TABLE V SAFETY INSTRUCTIONSRISK AND SAFETY PHRASES					
Buffer	Hazard Contents	Hazard Symbol		Risk Phrases	Safety Phrases
NT	Guanidine Thiocyanate	X Xn*	Harmful by inhalation, in contact with skin and if swallowed	R 20/21/22	S 13

Risk Phrases

R 20/21/22 Harmful by inhalation, in contact with skin and if swallowed

Safety Phrases

S 13 Keep away from food, drink and animal feedstuffs

^{*}Label not necessary, if quantity below 125 g or ml (concerning 67/548/EEC Art. 25, 1999/45/EC Art. 12 and German GefStoffV § 42 and TRGS 200 7.1)

V. Protocol for DNA Extraction from Agarose Gels

1. Excise DNA fragment

Take a clean scalpel to excise the DNA fragment from agarose gel. Excise gel slice containing the fragment carefully to minimize the gel volume. Determine the weight of the gel slice and transfer it to a clean tube.

2. Gel lysis

For each **100 mg** agarose gel add **200** μ *l* buffer NT.

For gels containing > 2% agarose, double the volume of buffer NT. The maximum amount of gel slice per NucleoSpin[®] Extract II column is 400 mg or 200 mg of a high percentage gel > 2%. In this case 2 loading steps are required (step 3).

Incubate sample at **50°C** until the gel slices are dissolved **(5-10 min)**. Vortex the sample briefly every 23 min until the gel slices are dissolved **completely!**

3. Bind DNA

Place a NucleoSpin[®] Extract II column into a 2 ml collecting tube and load the sample.

Centrifuge for **1 min at 11,000 x g.** Discard flow-through and place the NucleoSpin[®] Extract II column back into the collecting tube.

4. Wash silica membrane

Add **600** μ *I* **buffer NT3.** Centrifuge for **1 min** at **11,000 x g.** Discard flow-through and place the NucleoSpin[®] Extract II column back into the collecting tube.

V. Protocol for DNA Extraction from Agarose Gels continued

5. Dry silica membrane

Centrifuge for **2 min** at **11,000 x g** to remove **buffer NT3** quantitatively. Make sure the spin column doesn't come in contact with the flowthrough while removing it from the centrifuge and the collecting tube.

Residual ethanol from buffer NT3 would inhibit subsequent reactions and has to be removed in this step. In addition to centrifugation, total removal can be achieved by incubation of NucleoSpin[®] Extract II columns for 2-5 min at 70°C prior to elution.

6. Elute DNA

Place the NucleoSpin[®] Extract II column into a **clean** 1.5 ml microcentrifuge tube. Add **15-50** *µl* **elution buffer NE**, incubate at **room temperature** for **1 min** to increase the yield of eluted DNA. Centrifuge for **1 min** at **11,000 x g**.

Yield of larger fragments (> 5-10 kb) can be increased by using prewarmed elution buffer (70°C): For elution, add prewarmed elution buffer and incubate at room temperature for 1 min before collecting eluate by centrifugation.

VI. Protocol for Direct Purification of PCR Products

1. Adjust DNA binding conditions

Mix **2** volumes of buffer NT with **1** volume of sample (e.g. 200 μ l and 100 μ l PCR reaction mix).

For sample volumes $< 50 \,\mu$ l adjust the volume of the reaction mix to 50 μ l using TE buffer (pH 7.5).

2. Bind DNA

Place a NucleoSpin[®] Extract II column into a 2 ml collecting tube and load the sample.

Centrifuge for **1 min** at **11,000 x g.** Discard flow-through and place the NucleoSpin[®] Extract II column back into the collecting tube.

3. Wash silica membrane

Add **600** μ *l* **buffer NT3.** Centrifuge for **1 min** at **11,000 x g.** Discard flow-through and place the NucleoSpin[®] Extract II column back into the collecting tube.

4. Dry silica membrane

Centrifuge for 2 min at 11,000 x g to remove buffer NT3 quantitatively. Make sure the spin column doesn't come in contact with the flow-through while removing it from the centrifuge and the collecting tube.

Residual ethanol from buffer NT3 would inhibit subsequent reactions and has to be removed in this step. In addition to centrigugation, total removal can be achieved by incubation of NucleoSpin[®] Extract II columns for 2-5 min at 70°C prior to elution.

5. Elute DNA

Place the NucleoSpin[®] Extract II column into a **clean** 1.5 ml microcentrifuge tube. Add **15-50** μ *l* elution buffer NE, incubate at room temperature for 1 min to increase he yield of eluted DNA. Centrifuge for 1 min at 11,000 x g.

Yield of larger fragments (> 5-10 kb) can be increased by using prewarmed elution buffer (70°C): For elution, add prewarmed elution buffer and incubate at room temperature f or 1 min before collecting

VII. Support Protocol for Concentration, Desalination, Removal of Enzymes

1. Adjust DNA binding conditions

Add **2 volumes** of **buffer NT** to **1 volume** of the DNA containing **sample** (e.g. 200 μ / NT and 100 μ / PCR reaction mix).

If your sample contains large amounts of detergents or other critical substances, double the volume of NT.

2. Bind DNA

Continue with **step 2** of the protocol for purification of PCR products (section 5).

Appendix A: Troubleshooting

Problem	Possible cause and suggestions			
Incomplete	 High amount of agarose Use doubled volumes of buffer NT for highly concentrated and/or LMP (low melting point) agarose gels. 			
lysis of agarose slices	 Time and temperature Check incubation temperature. Depending on the weight of gel slice, incubation (section 4, step 2) can be prolonged up to 20 min. Vortex every 2 min and check integrity of the gel slice. Heavy weight gel slices may be quenched or crushed before addition of buffer NT. 			
	 Reagents not applied properly Add indicated volume of 96-100% ethanol to buffer NT3 concentrate and mix well before use. 			
	 Incompletely dissolved gel slice Increase time or add another two volumes of NT and vortex the tube every 2 minutes during incubation at 50°C. Small pieces of gel are hardly visible and contain DNA that will be lost for purification. 			
Low DNA yield	Insufficient drying of the NucleoSpin[®] Extract II silica membrane			
	 Centrifuge 5 min at 11,000 x g before elution to remove ethanolic buffer NT3 completely. Ethanolic contaminations are also indicated by gel-loading problems (samples float out of gel slots). Remove the spin cup carefully from centrifuge and and collecting tube and avoid contact of spin cup with flow-through. 			
	 Not enough elution buffer Especially when larger amounts of DNA (>5) are bound, increase elution buffer volume up to 100 μl. 			
	 Isolation of large DNA fragments Preheat elution buffer NE to 70°C and incubate on the 			

• Preheat elution buffer NE to 70°C and incubate on the silica membrane at room temperature for 2 min before centrifugation.

Appendix A: Troubleshooting continued

Problem Possible cause and suggestions

Carry-over of ethanol/ethanolic buffer NT3

 Be sure to centrifuge 3-5 min at 11,000 x g in order to achieve complete removal of ethanolic buffer NT3 after the washing step. Ethanolic contaminations are also indicated by gel-loading problems (samples float out of gel slots).

Suboptimal performance of DNA in sequencing reactions

Elution of DNA with Buffers other than buffer NE e.g. TE buffer (Tris/EDTA)

• EDTA may inhibit sequencing reactions. In this case it is recommended to repurify DNA and elute in buffer NE or water.

Not enough DNA used for sequencing reaction

 Quantitate DNA by agarose gel electrophoresis before setting up sequencing reactions.

Appendix B: Ordering Information

For a complete listing of all Clontech products, please visit **www.clontech.com**

<u>Product</u>	Pack of	Cat. No.
NucleoSpin [®] Extract II	10 preps	636971
NucleoSpin [®] Extract II	50 preps	636972
NucleoSpin [®] Extract II	250 preps	636973
NucleoSpin [®] collection tubes	1000	636029

Appendix C: References

Vogelstein B., and D. Gillespie. 1979. Preparative and analytical purification of DNA from agarose. Proc. Natl. Acad. Sci. USA **76:** 615-619

Appendix D: Product Use Restriction / Warranty

NucleoSpin® Extract II kit components were developed, designed and sold **for research purposes** only. They are suitable **for in vitro uses only.** No claim or representation is intended for its use to identify any specific organism or for clinical use (diagnostic, prognostic, therapeutic, or blood banking).

It is rather the responsibility of the user to verify the use of the **NucleoSpin® Extract II** kit for a specific application range as the performance characteristic of this kit has not been verified to a specific organism.

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Appendix D: Product Use Restriction / Warranty continued

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Notes

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