



Clean-up of Sequencing Reactions

User Manual

NucleoSEQ®



Plesmanlaan 1d 2333 BZ Leiden The Netherlands T. +31 (0)71 568 10 00 T. Belgium: 0800 71640 F. +31 (0)71 568 10 10 info@bioke.com www.bioke.com

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1 Components

1.1 Kit contents

	NucleoSEQ®			
	10 preps	50 preps	250 preps	
REF	740523.10	740523.50	740523.250	
NucleoSEQ® Columns	10	50	250	
Collection Tubes	10	50	250	
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1.2 Material to be supplied by user

• Collection tubes (e.g., 1.5 mL microcentrifuge tubes)

2 Product description

2.1 The basic principle

NucleoSEQ® Columns are designed for fast and effective clean-up of nucleic acids. Using gel exclusion in a convenient spin column format allows reliable removal of smaller molecules from nucleic acids. Impurities, for example, salts, excess of labels, nucleotides, traces of organic solvents, primers are retained by the column while nucleic acids of interest are recovered with high yield. The columns are pre-filled with size exclusion matrix.

2.2 Kit specifications

- Maximum sample volume to be loaded onto the column: 20 μL
- Removal of sequencing dye terminators including BigDye®

3 Storage conditions

- NucleoSEQ® Columns with dry gel matrix can be stored at room temperature for 12 months.
- NucleoSEQ® Columns with hydrated gel matrix should be stored at 4°C.
 Columns can be stored up to 14 days at 4°C.

4 Safety instructions – risk and safety phrases

The components of the NucleoSEQ® kits do not contain hazardous contents.

5 Protocol for sequencing reaction clean-up

General procedure

(For details on each step see page 7.)

1 Spin down dried gel resin



750 x *g* 30 s

2 Hydrate gel resin with 600 μL water, vortex, and incubate at least 30 min for complete hydration



600 µL water

RT > 30 min

3 Remove bottom plug and spin down hydrated gel resin



750 x *g* 2 min

4 Load sample to the center of the column



Load column

5 Spin for **4–6 min** at **750 x g** to recover purified sample



750 x *g* 4 – 6 min



Detailed procedure

Perform sequencing reaction according to standard protocols.

We recommend to use not more than $1-2~\mu L$ of Big Dye® Ready reaction mix in a 20 μL sequencing reaction in order to avoid overloading the column.

- 1 Centrifuge the NucleoSEQ® Columns for 30 s at 750 x g to collect the dry gel matrix on the bottom of the cartridge.
- 2 Add 600 μL dist. water and vortex to hydrate the gel matrix. Remove air bubbles by vortexing or tapping the column. Incubate at least 30 min or overnight to hydrate the gel matrix. Incubation can be performed at room temperature or 4°C. Hydrated columns can be stored at 4°C for a maximum of 14 days. Resuspend the settled gel matrix by inverting or vortexing the spin column several times. Air bubbles should not be visible now. Remove the bottom plug and place the spin column into a Collection Tube (supplied with the kit).
- 3 Place the column into an appropriate centrifuge (the hinge of the spin column's cover lid should be orientated to the outside of the rotor). Centrifuge 2 min at 750 x g to remove the remaining storage buffer. Discard the collection tube with storage buffer. Place the spin column in an appropriate collection tube (e.g., microcentrifuge tube, not supplied with the kit).
- 4 Open the lid of the column. Carefully load the sample drop by drop onto the center of the gel resin. Pipetting the sample at the sides of the spin column tube may reduce purification efficiency of the column. Moreover, do not disturb the gel surface. Sample volume should not exceed 20 μL.
- 5 Place the column in the same orientation as in step 3 into the centrifuge. Elute the sample by centrifuging the column for 4-6 min at 750 x g. Discard the spin column.

Dry the sample or use the sample directly.

6 Appendix

6.1 Troubleshooting

Problem	Possible cause and suggestions			
	Sample added to column improperly			
Dye blobs	 Add sample directly to the center of the settled gel matrix. Dispense sample dropwise. Avoid adding sample at the sides of the gel matrix 			
	Sample volume to high			
	- Add sample in a volume of 20 $\mu L.$ Higher sample volumes can cause incomplete removal of dye terminators.			
	Sample volume to small			
Poor signal intensity	• Add sample in a volume of 20 $\mu L.$ If necessary adjust sample volume to 20 μL using distilled water			

Conversion of RCF from different centrifuges:

 $rpm = 100 \text{ x } \sqrt{\frac{RCF}{1.12}} r$

 $RCF = (rpm/1000)^2 \times 1.12r$

rpm = revolutions per minute

RCF = relative centrifugal force (g force = RCF x g)

r = radius in mm

6.2 Ordering information

Product	REF	Pack of
NucleoSEQ®	740523.10	10 preps
	740523.50	50 preps
	740523.250	250 preps

Visit www.mn-net.com for more detailed product information.

6.3 Product use restriction/warranty

NucleoSEQ® kit components are intended, developed, designed, and sold FOR RESEARCH PURPOSES ONLY, except, however, any other function of the product being expressly described in original MACHEREY-NAGEL product leaflets.

MACHEREY-NAGEL products are intended for GENERAL LABORATORY USE ONLY! MACHEREY-NAGEL products are suited for QUALIFIED PERSONNEL ONLY! MACHEREY-NAGEL products shall in any event only be used wearing adequate PROTECTIVE CLOTHING. For detailed information please refer to the respective Material Safety Data Sheet of the product! MACHEREY-NAGEL products shall exclusively be used in an ADEQUATE TEST ENVIRONMENT. MACHEREY-NAGEL does not assume any responsibility for damages due to improper application of our products in other fields of application. Application on the human body is STRICTLY FORBIDDEN. The respective user is liable for any and all damages resulting from such application.

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IF THERE IS NO IVD SIGN, THE PRODUCT SHALL NOT BE SUITABLE FOR IN-VITRO-DIAGNOSTIC USE!

ALL OTHER PRODUCTS NOT LABELED AS IVD ARE NOT SUITED FOR ANY CLINICAL USE (INCLUDING, BUT NOT LIMITED TO DIAGNOSTIC, THERAPEUTIC AND/OR PROGNOSTIC USE).

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Please contact:

MACHEREY-NAGEL GmbH & Co. KG

Tel.: +49 (0) 24 21 969 270 e-mail: tech-bio@mn-net.com

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