

NucleoSpin<sup>®</sup> Microbial DNA

December 2022/Rev. 06



MACHEREY-NAGEL

www.mn-net.com

Bioanalysis

# Genomic DNA from microorganisms

## Protocol at a glance (Rev. 06)

1 Prepare sample		< 40 mg microbial pellet (wet weight)		
	V		Transfer sa	mole in
			MN Tube	Туре В
			40 μL Buf	fer MG
2 Lyse sample		Ò	10 μL Liquid P	roteinase K
	P		Agitate on a swing mi 4–12 r	ll or similar device nin
			11,000 x g	<i>g,</i> 30 s
			600 μL But	ffer MG
3 Adjust binding conditions		Ò	Vortex 3 s	
			11,000 x <i>g,</i> 30 s	
4 Rind DNA			Load 500–600 µL sample on NucleoSpin® Microbial DNA Column	
	ी॰व	<u> </u>	11,000 x g	<i>g,</i> 30 s
5 Wash silica		Ø	1st 500 μL BW	11,000 x <i>g,</i> 30 s
membrane			<b>2<sup>nd</sup></b> 500 μL B5	11,000 x <i>g</i> , 30 s
6 Drv silica	Ĩ			
membrane			11,000 x <i>g,</i> 30 s	
			100 μL Elutior	Buffer BE
7 Elute DNA		Ò	RT, 1 r	nin
			11,000 x g	<i>g,</i> 30 s

#### NucleoSpin® Microbial DNA



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# 1 Components

## 1.1 Kit contents

	Nucle	oSpin <sup>®</sup> Microbia	al DNA
REF	10 preps 740235.10	50 preps 740235.50	250 preps 740235.250
Lysis Buffer MG	10 mL	38 mL	5 x 38 mL
Wash Buffer BW	6 mL	30 mL	150 mL
Wash Buffer B5 (Concentrate)*	6 mL	6 mL	50 mL
Elution Buffer BE**	13 mL	30 mL	125 mL
Liquid Proteinase K	120 µL	600 µL	2 x 1.5 mL
MN Bead Tubes Type B	10	50	250
NucleoSpin <sup>®</sup> Microbial DNA Columns (light green rings)	10	50	250
Collection Tubes (2 mL)	20	100	500
User manual	1	1	1

 $<sup>^{\</sup>ast}\,$  For preparation of working solutions and storage, see section 3.

<sup>\*\*</sup> Composition of Elution Buffer BE: 5 mM Tris/HCl, pH 8.5

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

Reagents

• 96–100 % ethanol

Consumables

- 1.5 mL or 2 mL microcentrifuge tubes for microbial sample sedimentation
- Disposable tips

Equipment

- Manual pipettors
- Centrifuge for microcentrifuge tubes
- Vortex mixer
- Sample disruption device: swing mill or similar device (e.g., Mixer Mill MM200, MM300, MM400 (Retsch<sup>®</sup>); FastPrep<sup>®</sup> System (MP-Biomedicals); Precellys<sup>®</sup> (Bertin Technologies); MagNA Lyser (Roche); TissueLyser (QIAGEN); Bullet Blender<sup>®</sup> (Next Advance); Mini-Beadbeater (Biospec Products); Speed Mill (Analytik Jena); Vortex Adapter for Vortex-Genie<sup>®</sup> 2 X (QIAGEN)
- Personal protection equipment (lab coat, gloves, goggles)

#### 1.3 About this user manual

It is strongly recommended for first time users to read the detailed protocol sections of the **NucleoSpin® Microbial DNA** kit before using this product. 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 online at *www.mn-net.com*.

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

# 2 Product description

## 2.1 The basic principle

The **NucleoSpin<sup>®</sup> Microbial DNA** kit is designed for efficient isolation of genomic DNA from microbial samples. DNA can be isolated from a wide variety of microorganisms such as gram negative, and gram positive bacteria as well as yeast, e.g., *Escherichia coli, Bacillus subtilis, Corynebacterium glutamicum, Saccharomyces cerevisiae.* Preparation of the collected samples containing the microbes of interest should be in pellet format.

Preliminary data also indicate the usability of the kit for DNA isolation from fungal mycelia, e.g., *Aspergillus nidulans*, from bacterial spore suspensions, e.g., *Geobacillus stearothermophilus*, and from plant pollen, e.g., honey bee pollen baskets. For optimal DNA yield, bead tubes different from the ones included in the kit might be required for such applications (see section 2.4).

Microbial samples such as gram positive bacteria, yeast, and spores can be difficult to lyse due to their strong complex cell wall structures. The NucleoSpin<sup>®</sup> Microbial DNA kit replaces enzymatic lysis by utilizing mechanical disruption of cell wall structures with the MN Bead Tubes. The MN Bead Tubes can be used in combination with many compatible disruptive devices (see section 2.4.1). High DNA yields can be obtained with the MN Bead Tubes from a large variety of sample types – enabling the procedure to be convenient, fast, and easy. Alternative bead types can be ordered separately for select sample types (see section 2.4.2 for recommendations).

## 2.2 Kit specifications

Kit specifications at a glance			
Parameter	NucleoSpin <sup>®</sup> Microbial DNA		
Technology	Silica membrane technology		
Format	Mini spin column		
Sample material	Microbial cell culture pellets of gram positive and gram negative bacteria, yeast		
Sample amount	Up to approx. 40 mg wet weight		
Typical yield	Varies by sample and disruption device. 5–25 µg DNA from approx. 30 mg wet weight microbial pellet can be obtained		
A <sub>260</sub> /A <sub>280</sub>	1.6–2.0		
Elution volume	100–200 µL		
Preparation time	35 min/6 preps		
Binding capacity	60 µg		
Use	For research use only		

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

Cells should be harvested from fresh microbial cultures by sedimentation via centrifugation. Supernatant should be removed by aspiration. Microbial cell pellets can be used fresh or stored at -20  $^\circ$ C to -80  $^\circ$ C before starting DNA isolation.

#### 2.4 Lysis and disruption of sample material

In order to obtain optimal yields of DNA from sample material, a complete disruption of the sample material is necessary. Sample disruption efficiency depends on the following parameters and can be achieved by following suggestions outlined in the subsequent sections.

### 2.4.1 Disruption device

The following devices are compatible with MN Bead Tubes. Please check whether MN Bead Tubes can be accommodated by the available tube adapters prior to starting the procedure.

- MN Bead Tube Holder in combination with the Vortex-Genie® 2 (recommended).
- Mixer Mill MM200, MM300, MM400 (Retsch®) (suitable).

If other disruption devices (section 1.2) are intended to be used, consider section 2.4.2 and WARNING note in section 2.4.3!

## 2.4.2 Type of Bead tube

Bead type, disruption time, and frequency/speed must be optimized for a given sample to obtain maximal DNA yield and quality.

#### Type of Bead tube

- MN Bead Tubes Type A (0.6–0.8 mm ceramic beads) Recommended for soil, sediment, and stool (included in NucleoSpin<sup>®</sup> Soil, see ordering information, section 6.2).
- MN Bead Tubes Type B (40–400 μm glass beads) Recommended for gram positive and negative bacteria (included in NucleoSpin<sup>®</sup> Microbial DNA, see ordering information, section 6.2).
- MN Bead Tubes Type C (1–3 mm corundum) Recommended for yeast (see ordering information, section 6.2).
- MN Bead Tubes Type D (3 mm steel beads; included in NucleoSpin<sup>®</sup> DNA Insect kits)
   Recommended for insects, crustaceans, and lipid-rich tissue.
- MN Bead Tubes Type E (combination of 3 mm steel beads and 40–400 µm glass beads) Recommended for hard to lyse bacteria within insect or tissue samples (see ordering information, section 6.2).
- MN Bead Tubes Type F (combination of 1-3 mm corundum and 3 mm steel beads;) Recommended for challenging tissues, e.g., spleen, or lung tissue.)
- MN Bead Tubes Type G (5 mm steel beads) Recommended for plant material (see ordering information, section 6.2).

## 2.4.3 Time and frequency of disruption

The following recommendations have been established for the MN Bead Tube Holder in combination with a Vortex-Genie<sup>®</sup> 2 or a Retsch<sup>®</sup> Mixer Mill MM300 operating at highest frequency (30 Hertz). For using other disruption devices, and other sample materials, time and frequency have to be optimized.

#### Time and frequency of disruption using MN Bead Tube Holder on a Vortex Genie® 2

As a general starting point disrupt microbial samples for 20 min using MN Bead Tube Holder on a Vortex Genie<sup>®</sup> 2.

#### Time an frequency of disruption using a Retsch® Mixer Mill MM300

Sample material	MN Bead Tube	Disruption time
Gram negative bacteria E.g., Escherichia coli, Vibrio fischeri	MN Bead Tubes Type B (Alternative: Type A, Type C)	4 min
<b>Gram positive bacteria</b> E.g., <i>Bacillus subtilis</i> , <i>Corynebacterium glutamicum</i>	MN Bead Tubes Type B (Alternative: Type A)	12 min
<b>Yeast</b> E.g., <i>Saccharomyces</i> <i>cerevisiae</i>	MN Bead Tubes Type C	12 min
<b>Filamentous fungi</b> E.g., Aspergillus spec., Rhizopus spec.	MN Bead Tubes Type C	12 min

<u>Note:</u> Performance and stability testing has been conducted on the MN Bead Tubes A, B, and C on a Retsch<sup>®</sup> Mixer Mill MM300 at highest frequency (30 Hertz) for up to 15 minutes for optimal sample disruption, avoidance of DNA fragmentation, and tube durability. Other disruption devices (see section 2.4.1) will require different settings regarding frequency and duration for optimal performance with the selected sample material. Please note that the position of the tube within the machine (Retsch<sup>®</sup> Mixer Mill) is important for optimal performance! Please consult instruction manual of the machine.

**WARNING:** Many modern disruption devices can cause very high energy input in bead tubes. Depending on bead tube type and content (beads, liquid volume, sample type), especially high frequency of shaking and / or long shaking duration can cause breaking up of the bead tubes! It is the responsibility of the user to perform initial stability test for the used bead tubes under the conditions used! Perform initial test with water instead of lysis buffer and moderate machine setting (low frequency, short time) in order to avoid spillage of chaotropic lysis buffer in case of tube breakage.

**WARNING:** In section 5 a certain liquid volume during disruption is recommended. The reduction of liquid will severely increase the mechanical impact of the grinding matrix and can result in damage of DNA and tube (especially if MN Bead Tubes D and E are used).

#### 2.5 Elution procedures

In addition to the standard method, several modifications are possible to increase yield, concentration, and convenience.

- **Convenient elution (standard elution)**: For convenience, elution can be performed by one time addition of 100 µL elution buffer onto the column.
- High yield: Two serial elutions of 100 μL each for total elution volume of 200 μL.
- High concentration: Use initial 100 µL eluate for second elution 100 µL total elution volume, 2 elutions.

For more information please visit our cultured microorganism website. *www.mn-net.com/culturedmicroorganism* 

# 3 Storage conditions and preparation of working solutions

#### Attention:

Lysis Buffer MG and Wash Buffer BW contain chaotropic salt! Wear gloves and goggles!

CAUTION: Buffers MG and BW contain chaotropic salts which can form highly reactive compounds when combines with bleach (sodium hypochlorite). DO NOT add bleach or acidic solutions directly to the sample-preparation waster!

 All kit components can be stored at 15–25 °C and are stable until: see package label.

Before starting any NucleoSpin® Microbial DNA protocol, prepare the following:

- Wash Buffer B5: Add the indicated volume of ethanol (96–100 %) to Wash Buffer B5 Concentrate. Mark the label of the bottle to indicate that ethanol was added. Wash Buffer B5 can be stored at 15–25 °C for at least one year.
- Liquid Proteinase K is ready to use. After first time use, store Liquid Proteinase K at 4 °C or -20 °C.

NucleoSpin <sup>®</sup> Microbial DNA					
REF	10 preps	50 preps	250 preps		
	740235.10	740235.50	740235.250		
Wash Buffer B5	6 mL	6 mL	50 mL		
(Concentrate)	Add 24 mL ethanol	Add 24 mL ethanol	Add 200 mL ethanol		

# 4 Safety instructions

When working with the **NucleoSpin<sup>®</sup> Microbial DNA** 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*).



Caution: Guanidine hydrochloride in Buffer BW and guanidinium thiocyanate in buffer MG 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® Microbial DNA** 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 and Proteinase K 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 Protocols

#### 5.1 Protocol for gram positive and gram negative bacteria

#### Before starting the preparation:

- Check if Buffer B5 was prepared according to section 3.
- Check section 2.4 for lysis and disruption of sample material.

#### 1 Prepare sample

	Harvest cells from a culture by centrifugation in a microcentrifuge tube (not provided). Discard supernatant. <i>Up to approximately 40 mg of wet weight microbial cell</i>	+ 100 μL BE
	culture pellet can be used as sample material.	
	Add 100 µL Elution Buffer BE and resuspend cells.	
	Alternatively, high quality grade water (not provided) can be used.	
2	Lyse sample	
	Transfer the cell suspension into the <b>MN Bead Tube Type B</b> (provided).	+ 40 μL MG + 10 μL Liquid
	Add 40 μL Buffer MG. Then, add 10 μL Liquid Proteinase K and close the tube.	Proteinase K
	Note: It is not necessary to vortex here.	
	Agitate the MN Bead Tube on a swing mill or similar device.	Agitate
	<u>Note:</u> Optimal agitation duration, speed/frequency depends on the machine used. On a Retsch <sup>®</sup> Mixer Mill MM200, MM300, MM400, e.g., 4 min at maximal frequency (30 Hertz) is adequate for E. coli, 12 min for B. subtilis (see section 2.4). On the swing mill, position of the tube in the mill can considerably influence the result. Please consult the instruction manual of the device used.	
	Centrifuge the MN Bead Tube <b>30 s</b> at <b>11,000 x</b> <i>g</i> to clean the lid.	💍 11,000 x g,

3	Adjust DNA binding conditions		
	Add $600~\mu L$ Buffer MG and mix (e.g, vortex for 3 s).		+ 600 μL MG
	<u>Note:</u> Glass beads should be resuspended; some residual pellet (cell debris) may remain on the bottom of the tube.		Mix
	Centrifuge for <b>30 s</b> at <b>11,000 x</b> <i>g</i> .	$\bigcirc$	30 s
	Note: This centrifugation step is performed in order to clean the lid and sediment glass beads and cell debris.		
4	Bind DNA		
	<b>Transfer</b> the supernatant (~500–600 µL) onto the <b>NucleoSpin<sup>®</sup> Microbial DNA Column</b> , placed in a 2 mL Collection Tube (provided).	jo mood	Load samples 11.000 x <i>a</i> .
	Centrifuge for <b>30 s</b> at <b>11,000 x g</b> . Discard collection tube with flowthrough. Put column into a fresh Collection Tube (2 mL, provided).	Ö	30 s
5	Wash silica membrane		
	1 <sup>st</sup> wash		+ 500 μL BW
	Add <b>500 µL Buffer BW</b> . Centrifuge for <b>30 s</b> at <b>11,000 x g</b> . Discard flowthrough and place the column back into the Collection Tube.		11,000 x <i>g</i> , 30 s
	2 <sup>nd</sup> wash	<b>9</b>	+ 500 ul B5
	Add <b>500 µL Buffer B5</b> to the column and centrifuge for <b>30 s</b> at <b>11,000 x</b> <i>g</i> . Discard flowthrough and place the		+ σου με Βο
	column back into the Collection Tube.	Ò	11,000 x <i>g</i> , 30 s
6	Dry silica membrane	$\sim$	44.000
	Centrifuge the column for <b>30 s</b> at <b>11,000 x</b> <i>g</i> .		11,000 x <i>g</i> , 30 s
	Note: Residual wash buffer is removed in this step.	Ŭ	

#### 7 Elute highly pure DNA

Place the NucleoSpin<sup>®</sup> Microbial DNA Column into a 1.5 mL nuclease-free tube (not provided) and add 100  $\mu$ L Elution Buffer BE onto the column. Incubate at room temperature for 1 min. Centrifuge 30 s at 11,000 x g. + 100 μL BE RT, 1 min 11,000 x g, 30 s

For alternative elution procedures see section 2.5.

#### 5.2 Protocol for yeast (e.g., Saccharomyces cerevisiae)

Optimal DNA yields from yeast samples can be obtained by following the standard protocol using MN Bead Tube Type C (see ordering information on section 6.2) instead of MN Bead Tube Type B provided with the NucleoSpin<sup>®</sup> Microbial DNA kit.

The agitation is recommended at a Retsch<sup>®</sup> Mixer Mill MM300: 12 min at 30 Hz. For other disruption devices, please check section 2.4. Please note that the position of the tube within the machine is important for optimal performance, please consult instruction manual of the machine.

If bead carryover is observed in the eluate, transfer the eluate into a new 1.5 mL nucleasefree tube carefully avoid disturbing the pellet.

# 6 Appendix

# 6.1 Troubleshooting

Problem	Possible cause and suggestions			
	Incomplete lysis			
	<ul> <li>Adjust lysis conditions (bead tube type, agitation device, duration, or frequency).</li> </ul>			
	Reagents not applied properly			
	• Prepare Buffer B5 according to instructions (section 3).			
	Suboptimal elution of DNA from the column			
No or poor DNA yield	<ul> <li>For certain sample types, preheat Buffer BE to 70 °C before elution. Apply Buffer BE directly onto the center of the silica membrane.</li> </ul>			
	<ul> <li>Elution efficiencies decrease dramatically, if elution is done with buffers with a pH &lt; 7.0. Use slightly alkaline elution buffers like Buffer BE (pH 8.5).</li> </ul>			
	<ul> <li>Especially when expecting high yields from large amounts of material, we recommend elution with 200 µL Buffer BE and incubation of the closed columns in an incubator at 70 °C for 5 min before centrifugation.</li> </ul>			
	High A <sub>260</sub> /A <sub>280</sub> ratio			
Poor DNA quality	• Ratios > 1.9 can be caused by RNA contamination. Usually, such RNA contamination do not interfere with downstream application. Depending on sample type, amount, and disruption procedure, preparations might contain small amounts of RNA. If it is necessary to reduce RNA contamination to the lowest possible level, incubate the lysate after the disruption step for 5 min at 70 °C in order to inactivate the Proteinase K. After cooling to room temperature, add 20 $\mu$ L RNase A (20 mg/mL) and incubate 5 min. Continue with the application of the lysate onto the column.			
	Reagents not applied properly			
	• Prepare Buffer B5 according to instructions (see section 3).			
	Too much sample material used			
Clogged columns	• Make sure to centrifuge the lysate after cell disruption in order to sediment beads and cell debris. Only transfer cleared supernatant onto the column.			

Problem	Possible cause and suggestions
	Carry-over of ethanol or salt
Suboptimal	• Make sure to centrifuge ≥ 1 min at 11,000 x g in order to remove all of ethanolic Buffer B5 before eluting the DNA. If, for any reason, the level of Buffer B5 has reached the column outlet after drying, repeat the centrifugation.
genomic DNA in enzymatic reactions	• Do not chill Buffer B5 before use. Cold buffer will not remove salt effectively. Equilibrate Buffer B5 to room temperature before use.
	Contamination of DNA with inhibitory substances

• Do not elute DNA with TE buffer. EDTA may inhibit enzymatic reactions. Repurify DNA and elute in Buffer BE.

## 6.2 Ordering information

Product	REF	Pack of
NucleoSpin <sup>®</sup> Microbial DNA	740235.10/.50/.250	10/50/250 preps
MN Bead Tube Holder	740469	1 piece
NucleoSpin <sup>®</sup> Soil	740780.10/.50/.250	10/50/250 preps
NucleoSpin <sup>®</sup> DNA Lipid Tissue	740471.10/.50	10/50 preps
NucleoSpin <sup>®</sup> DNA Insect	740470.10/.50	10/50 preps
NucleoSpin <sup>®</sup> DNA Stool	740472.10/.50	10/50 preps
MN Bead Tubes Type A (0.6–0.8 mm ceramic beads, recommended for soil and sediments)	740786.50	50 pieces
MN Bead Tubes Type B (40–400 $\mu m$ glass beads, recommended for bacteria)	740812.50	50 pieces
MN Bead Tubes Type C (1–3 mm corundum, recommended for yeast)	740813.50	50 pieces
MN Bead Tubes Type D (3 mm steel beads, recommended for insects)	740814.50	50 pieces
MN Bead Tubes Type E (40-400 µm glass beads and 3 mm steel beads, recommended for hard to lyse bacteria within insect samples)	740815.50	50 pieces

Product	REF	Pack of
MN Bead Tubes Type F (1-3 mm corundum and 3 mm steel beads, recommended for challenging tissues, e.g., spleen, or lung tissue)	740816.50	50 pieces
MN Bead Tubes Type G (5 mm steel beads, recommended for plant material)	740817.50	50 pieces
Buffer BE	740306.100	125 mL
Buffer B5 Concentrate (for 125 mL Buffer B5)	740921	25 mL
Buffer BW	740922	100 mL
Liquid Proteinase K	740396	5 mL
RNase A	740505.50 740505	50 mg 100 mg
Collection Tubes (2 mL)	740600	1000

#### 6.3 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-NAGELS 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 costumer and are not a part of a contract of sale or of this warranty.

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Products and their application are subject to change. Therefore, please contact our Technical Service Team for the latest information on MACHEREY-NAGEL products. You may also contact your local distributor for general scientific information. Descriptions in MACHEREY-NAGEL literature are provided for informational purposes only.

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Plasmid DNA Clean up RNA DNA Viral RNA and DNA Protein High throughput Accessories Auxiliary tools



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