

NucleoMag[®] DNA FFPE

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MACHEREY-NAGEL

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Bioanalysis

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1 Components

1.1 Kit contents

	NucleoMag [®] DNA FFPE		
REF	1 × 96 preps 744320.1	4 × 96 preps 744320.4	
NucleoMag [®] B-Beads	1.8 mL	4 × 1.8 mL	
Lysis Buffer FL	30 mL	4 × 30 mL	
Binding Buffer MB2	100 mL	500 mL	
Wash Buffer MB4	75 mL	300 mL	
Elution Buffer MB6	30 mL	125 mL	
Proteinase K (lyophilized)*	75 mg	4 × 75 mg	
Proteinase Buffer PB	15 mL	35 mL	
Paraffin Dissolver (blue)	60 mL	2 × 125 mL	
Decrosslink Buffer D-Link	30 mL	2 × 30 mL	
User manual	1	1	

^{*} For preparation of working solutions and storage conditions see section 3.

1.2 Reagents, equipment and consumables to be supplied by user

Reagents

• 80 % ethanol

Equipment/Consumables

Product	REF	Pack of
Magnetic separation system e.g., NucleoMag [®] SEP (see section 2.4)	744900	1
Separation plate for magnetic beads separation, e.g., Square-well Block (96-well block with 2.1 mL square- wells)	740481 740481.24	4 24
Lysis tubes for incubation of samples and lysis, e.g., Rack of Tubes Strips (1 set consists of 1 Rack, 12 Strips with 8 tubes (1.2 mL wells) each, and 12 Cap Strips)	740477 740477.24	4 sets 24 sets
Elution plate for collecting purified nucleic acids, e.g., Elution Plate U-bottom (96-well 0.3 mL microtiterplate with 300 µL u-bottom wells) e.g., Elution Plate Flat-bottom (96-well 0.3 mL microtiterplate with 300 µL flat-bottom wells)	740486 740673	24 20
For use of kit on KingFisher [®] 96 instrument: e.g., KingFisher [®] 96 Accessory Kit A (Square-well Blocks, Deep-well Tip Combs, Elution Plates for 4 × 96 NucleoMag [®] DNA FFPE preps using KingFisher [®] 96 platform)	744950	1 set

1.3 About this user Manual

It is recommended to read the instructions of this user manual carefully before use. All technical literature is also available on the internet at **www.mn-net.com**.

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

2 Product description

Formalin-fixed, paraffin-embedded (FFPE) tissue samples are routinely prepared from human surgical tissue samples by fixation with formalin and embedding in paraffin. Thin sections of FFPE samples are commonly subjected to histopathological analysis and remaining paraffin-tissue blocks are usually archived. Existing extensive archives of FFPE tissue samples represent a valuable source for retrospective studies of gene expression patterns and mutation analysis. However, the use of such samples for DNA analysis is limited due to chemical modification by formaldehyde and fragmentation of the DNA during tissue processing (sampling, fixing, embedding) and storage (humidity, time, temperature) of the samples. Standard DNA isolation procedures often result in low DNA yield or poor performance in downstream applications (e.g., PCR).

2.1 The basic principle

The **NucleoMag® DNA FFPE** kit is designed for the isolation of DNA from formalin-fixed, paraffin-embedded (FFPE) tissue specimen. The procedure replaces the use of flammable and malodorous xylene or d-limonene commonly used for deparaffinization. Further, the procedure does not require the difficult removal of organic solvent from often barely visible tissue pellets. **NucleoMag® DNA FFPE** employs the odorless Paraffin Dissolver (patent pending) and allows effective lysis in a convenient two-phase system. This kit provides reagents and magnetic beads for a convenient, reliable, and fast method to isolate DNA from 96 or 384 samples. First, the paraffin of FFPE sections is dissolved in the Paraffin Dissolver. Tissue is then digested by proteinase to solubilize the fixed tissue and release DNA into solution. Subsequently, heat incubation with a specially formulated buffer effectively eliminates crosslinks from the previously released DNA. For binding of the DNA to the paramagnetic beads, Binding Buffer MB2 and the NucleoMag® B-Beads are added to the lysate.

The procedure is based on the reversible adsorption of nucleic acids to paramagnetic beads under appropriate buffer conditions. After magnetic separation, the paramagnetic beads are washed to remove contaminants and salts using Wash Buffers MB4 and 80 % ethanol. Residual ethanol from previous wash steps is removed by airdrying. Finally, highly pure DNA is eluted with low salt Elution Buffer MB6. Purified DNA can directly be used for downstream applications, e.g., qPCR. The **NucleoMag® DNA FFPE** kit can be used either manually or automated on standard liquid handling instruments or automated magnetic separators.

2.2 Kit specifications

The **NucleoMag® DNA FFPE** kit is recommended for the isolation of DNA from formalinfixed, paraffin-embedded (FFPE) tissue samples. Samples are typically thin sections (approx. 3–20 µm thickness) of human or animal origin usually obtained by tissue resection or biopsy.

- Sample amount: The maximum sample size is determined by a) the amount of tissue and b) by the amount of paraffin. NucleoMag[®] DNA FFPE is suitable for up to 5 mg tissue. The amount of paraffin is limited to 15 mg, when using the standard protocol with Paraffin Dissolver (approx. 7 sections of 10 µm x 250 mm²). However, larger amounts of paraffin samples may be processed by using either additional Paraffin Dissolver or by deparaffinization using xylene.
- DNA yield strongly depends on the sample type, quality, quantity as well as conditions and duration of storage. Further, measured DNA yield may vary considerably between different quantification methods. Yield determined by absorption measurement at 260 nm or by a fluorescent dye (e.g., PicoGreen[®]) may deviate from values obtained by

quantification with PCR. Even quantification values obtained via PCR with a short (e.g., 80 bp) and a long (e.g., 300 bp) amplicon may also differ considerably. The deviation of quantification also depends on DNA size distribution as well as on efficiency of decrosslinking (or extent of remaining crosslinks).

- DNA size distribution: DNA isolated from formalin-fixed, paraffin-embedded tissue shows size distribution from 50 to 5,000 bases. Predominantly DNA of approx. 100–300 bases is observed, especially when the sample material is old. However, samples which were subjected to good tissue fixation, embedding, and storage conditions can yield DNA even larger than 5,000 bp.
- DNA preparation time strongly depends on the sample and the required lysis time. For best results lysis is performed at room temperature for at least three hours. For some kinds of samples a longer lysis (e.g., overnight) will even result in remarkably higher DNA yield.
- For research use only

NucleoMag[®] DNA FFPE is designed for use with NucleoMag[®] SEP magnetic separator plate (see ordering information, section 6.2) or other magnetic separation systems (see section 2.4). Manual preparation time of 96 samples is about 120 minutes.

NucleoMag[®] DNA FFPE allows easy automation on common liquid handling instruments or automated magnetic separators. The actual processing time depends on the configuration of the instrument and the magnetic separation system used. Typically, 96 samples can be purified in less than 120 minutes using the NucleoMag[®] SEP on the automation platform.

2.3 Handling, preparation, and storage of starting materials

Many factors influence the yield and quality of DNA obtained from FFPE samples. The procedure of tissue sampling, post sampling delay before fixation, fixation time, embedding, and storage conditions have a high impact on DNA quality and yield. Starting from a paraffin embedded tissue block, samples should be sectioned under clean conditions. Paraffin sections may be stored at +4 °C or lower for at least several weeks without observable effects on DNA yield or usability. Long term storage of paraffin sections may have a negative effect on the DNA due to air oxidation. Wear gloves at all times during the preparation. Change gloves frequently.

2.4 Magnetic separation systems

For use of **NucleoMag[®] DNA FFPE**, the use of the magnetic separator NucleoMag[®] SEP is recommended. Separation is carried out in a Square-well Block (see ordering information, section 6.2). The kit can also be used with other common separators

Magnetic separator	Separation plate or tube
NucleoMag [®] SEP (MN REF 744900)	Square-well Block (MN REF 740481 / .24)
Tecan Te-MagS™	1.5 mL tubes without lid (Sarstedt)

Static magnetic pins

Separators with static magnetic pins, for example, NucleoMag[®] SEP (for manual use and for use on liquid handling workstations): This type of separator is recommended in combination with a suitable microplate shaker for optimal resuspension of the beads during the washing and elution steps. Alternatively, beads can be resuspended in the buffer by pipetting up and down several times. For fully-automated use on liquid handling workstations, a gripper tool is required, the plate is transferred to the magnetic separator for separation of the beads and transferred to the shaker module for resuspension of the beads.

Movable magnetic systems

Separators with moving magnetic pins: Magnetic pins/rods are moved from one side of the well to the other and vice versa. Beads follow this movement and are thus pulled through the buffer during the wash and elution steps. Separation takes place when the system stops.

Automated separators

Separators with moving magnets: Magnetic beads are transferred into suitable plates or tubes. Beads are resuspended from the rod-covered magnets. Following binding, washing or elution beads are collected again with the rod-covered magnets and transferred to the next plate or tube.

2.5 Adjusting the shaker settings

When using a plate shaker for the washing and elution steps, the speed settings have to be adjusted carefully for each specific separation plate and shaker to prevent cross-contamination from well to well. Proceed as follows:

Adjusting shaker speed for binding and wash steps:

- Load 600 µL dyed water to the wells of the separation plate. Place the plate on the shaker and start shaking with a moderate speed setting for 30 seconds. Turn off the shaker and check the plate surface for small droplets of dyed water.
- Increase speed setting, shake for an additional 30 seconds, and check the plate surface for droplets again.
- Continue increasing the speed setting until you observe droplets on top of the separation plate. Reduce speed setting, check again, and use this setting for the washing step.

Adjusting shaker speed for the elution step:

 Load 100 µL dyed water to the wells of the collection plate and proceed as described above.

2.6 Handling of beads

Distribution of beads

A homogeneous distribution of the magnetic beads to the individual wells of the separation plate is essential for a high well-to-well consistency. Therefore, before distributing the beads, make sure that the beads are completely resuspended. Shake the storage bottle well or place it on a vortexer shortly. Premixing magnetic beads with the binding buffer allows easier homogenous distribution of the beads to the individual wells of the separation plate. During automation, a premix step before aspirating the beads / binding buffer mixture from the reservoir is recommended to keep the beads resuspended.

Magnetic separation time

Attraction of the magnetic beads to the magnetic pins depends on the magnetic strength of the magnetic pins, the selected separation plate, distance of the separation plate from the magnetic pins, and the volume to be processed. The individual times for complete attraction of the beads to the magnetic pins should be checked and adjusted on each system. It is recommended using the separation plates or tubes specified by the supplier of the magnetic separator.

Washing the beads

Washing the beads can be achieved by shaking or mixing. In contrast to mixing by pipetting up and down, mixing by shaker or magnetic mixing allows simultaneous mixing of all samples. This reduces the time and number of tips needed for the preparation. Resuspension by pipetting up and down, however, is more efficient than mixing by a shaker or magnetic mix.

Method	Resuspension efficiency	Speed	Number of tips needed
Magnetic mix	+	++	Low
Shaker	++	++	Low
Pipetting	+++	+*	High

+: acceptable, ++: good, +++: excellent, * 8-channel pipetting device

2.7 Handling of Proteinase K

For dispensing the Proteinase K solution to each sample it is recommended to predispense the needed amount to a separate reaction tube. Using a liquid handling device it is recommended to dispense the needed Proteinase K solution (25 μ L per prep) with 10% extra volume in a suitable tube for the correspondent robot. Unused Proteinase K solution should be stored at - 20 °C for further extractions.

2.8 Elution procedures

Purified DNA can be eluted directly with the supplied Elution Buffer MB6. Elution can be carried out in a volume of $\geq 25 \ \mu$ L. It is essential to cover the NucleoMag[®] Beads completely with elution buffer during the elution step. The volume of dispensed elution buffer depends on the magnetic separation system (e.g., the position of the pellet inside the separation plate). For efficient elution, the magnetic bead pellet should be resuspended completely in the elution buffer. For some separators, higher elution volumes might be necessary to cover the whole pellet.

Elution is possible at room temperature. Yield can be increased by 15-20 % if elution is performed at 55 °C.

3 Storage conditions and preparation of working solutions

Attention: Buffers MB2 and MB4 contain chaotropic salt! Wear gloves and goggles!

Storage conditions:

- All components of the NucleoMag[®] DNA FFPE kit should be stored at room temperature (15-25 °C) and are stable until: see package label.
- All buffers are delivered ready to use.

Before starting any NucleoMag[®] DNA FFPE protocol, prepare the following:

Proteinase K: Before first use of the kit, add the indicated volume of Proteinase Buffer PB to dissolve lyophilized Proteinase K. Proteinase K solution is stable at - 20 °C for at least 6 months.

NucleoMag [®] DNA FFPE		
REF	1 × 96 preps 744320.1	4 × 96 preps 744320.4
Proteinase K (lyophilized)	75 mg	75 mg
	Add 2.8 mL Proteinase Buffer	Add 2.8 mL Proteinase Buffer

• 80% ethanol: Use molecular biology grade ethanol, dilute with appropriate nuclease-free water to 80%.

4 Safety instructions

When working with the **NucleoMag[®] DNA FFPE** 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: Sodium perchlorate in buffer MB2 and buffer MB4 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 **NucleoMag® DNA FFPE** 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 Protocol for the isolation of genomic DNA from tissue

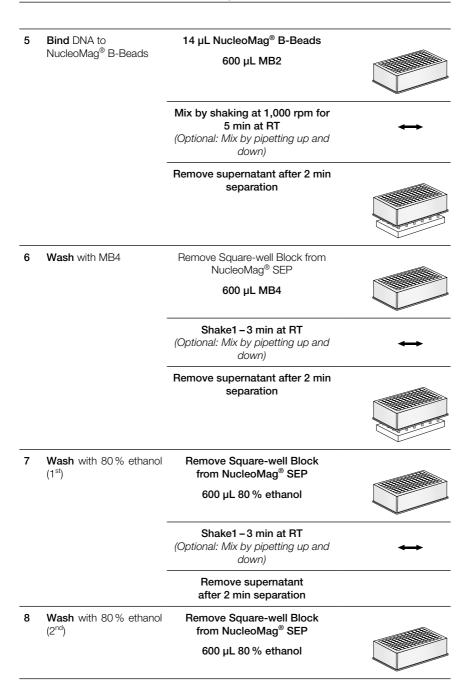
Protocol at a glance

- For additional equipment and hardware requirements, refer to section 1.2 and 2.4, respectively.
- For detailed information on each step, see page 16.

Before starting the preparation:

• Check if Proteinase K was prepared according to section 3.

1	Deparaffinize sample	400 μL Paraffin Dissolver 60 °C, 3 min	
		Mix hot sample by vortexing	
		Cool down to RT	0
2	Lyse sample	Add to each sample: 200 μL FL 25 μL Proteinase K	
		Mix by vortexing for 5 s 11,000 x <i>g</i> , 1 min 56 °C, 1 – 3 h	Õ
3	Decrosslink sample	11,000 x <i>g</i> , 30 s	
		100 μL D-Link	Ó
		Mix by vortexing for 5 s	
		11,000 x <i>g</i> , 1 min	V
		90 °C , 30 min	Ċ
		11,000 x <i>g</i> , 1 min	
4	Transfer sample	Load aqueous phase (lower) in Square-well Block	



		Shake1 – 3 min at RT (Optional: Mix by pipetting up and down)	↔
		Remove supernatant after 2 min separation	
9	Air dry magnetic beads	Air dry for 10 min at RT	
10	Elute DNA	Remove Square-well Block from NucleoMag [®] SEP	
		25-100 μL MB6 (Optional: Elute at 56 °C)	
		Shake 5 min at RT (Optional: Mix by pipetting up and down)	↔
		Separate 2 min and transfer DNA into elution plate / tubes	

Detailed protocol

This protocol is designed for magnetic separators with static pins (e.g., NucleoMag[®] SEP) and suitable plate shakers (see section 2.4). It is recommended using a Square-well Block for separation (see section 1.2). Alternatively, isolation of DNA can be performed in reaction tubes with suitable magnetic separators. This protocol is for manual use and serves as a guideline for adapting the kit to robotic instruments.

Before starting the preparation:

• Check if Proteinase K was prepared according to section 3.

1 Deparaffinize sample

Place the sample into a suitable 1.5 mL microreaction tube.

Add **400 \muL Paraffin Dissolver** to the sample. Incubate **3 min** at **60 °C** (to melt the paraffin). **Vortex or shake** the sample immediately (at 60 °C) at a vigorous speed to dissolve the paraffin. Cool down sample to room temperature.

Make sure that paraffin completely melts during the heat incubation step and mix well after melting to completely dissolve the paraffin.

Insufficient mixing of the heated sample may cause recurrence of solid paraffin particles. Make sure the sample does not comprise more than 15 mg paraffin or adjust the volume of Paraffin Dissolver.

2 Lyse sample

Add 200 μ L Lysis Buffer FL and 25 μ L Proteinase K to the lower aqueous phase. Mix well by repeated pipetting up and down, or pulse vortexing, or shaking. Centrifuge at 11,000 x g for 1 min and incubate at 56 °C for 1 – 3 h or overnight with shaking.

3 Decrosslink sample

Centrifuge for 1 min at 11000 x g.

Set heating block to 90 °C. Add **100 \muL Buffer D-Link**, mix by repeated pipetting up and down, vortexing for 5 s, or shaking.

Centrifuge for 1 min at 11,000 x g to obtain phase formation. Incubate at 90 °C for 30 min, mix by vortexing for 5 s. Cool down samples to room temperature

Centrifuge samples for 1 min at 11,000 x g.

Decrosslinking step is strongly recommended for short time lysis (1-3 h) and may be omitted after overnight lysis.

4 Load sample to Square-well Block

Transfer $400 \ \mu L$ of the lower aqueous phase from each sample to a Square-well Block for further processing.

5 Bind DNA to NucleoMag® B-Beads

Add 14 µL of NucleoMag[®] B-Beads and 600 µL Buffer MB2 to the lysed sample.

Mix by pipetting up and down 6 times and shake for 5 min at room temperature.

Alternatively, when processing the kit without a shaker, pipette up and down 10 times and incubate for 5 min at room temperature.

<u>Note:</u> Be sure to resuspend the NucleoMag[®] B-Beads before removing them from the storage bottle. Vortex storage bottle briefly until a homogenous suspension has been formed.

Separate the magnetic beads against the side of the wells by placing the Squarewell Block on the NucleoMag[®] SEP a magnetic separator. Wait at least 2 min until all the beads have been attracted to the magnets. Remove and discard supernatant by pipetting.

Note: Do not disturb the attracted beads while aspirating the supernatant.

6 Wash with MB4

Remove the Square-well Block from the NucleoMag[®] SEP magnetic separator.

Add **600 µL Buffer MB4** to each well and mix the beads by shaking until the beads are resuspended completely (**1 – 3 min**). Alternatively, resuspend beads completely by repeated pipetting up and down (15 times).

Separate the magnetic beads by placing the Square-well Block on the NucleoMag[®] SEP magnetic separator. Wait at least **2 min** until all the beads have been attracted to the magnet. Remove and discard supernatant by pipetting.

7 Wash with 80 % ethanol (1st)

Remove the Square-well Block from the NucleoMag[®] SEP magnetic separator.

Add **600 µL 80 % ethanol** to each well and mix the beads by shaking until the beads are resuspended completely (**1 – 3 min**). Alternatively, resuspend beads completely by repeated pipetting up and down (15 times).

Separate the magnetic beads by placing the Square-well Block on the NucleoMag[®] SEP magnetic separator. Wait at least **2 min** until all the beads have been attracted to the magnet. Remove and discard supernatant by pipetting.

8 Wash with 80 % ethanol (2nd)

Remove the Square-well Block from the NucleoMag® SEP magnetic separator.

Add 600 μ L 80 % ethanol to each well and resuspend the beads by shaking until the beads are resuspended completely (1 – 3 min). Alternatively, resuspend beads completely by repeated pipetting up and down (15 times).

Separate the magnetic beads by placing the Square-well Block on the NucleoMag[®] SEP magnetic separator. Wait at least **2 min** until all the beads have been attracted to the magnet. Remove and discard supernatant by pipetting.

9 Air dry

Air dry the magnetic bead pellet for **10 min** at **room temperature**.

10 Elution

Add desired volume of **Buffer MB6 (25–100 \muL)** to each well of the Square-well Block and resuspend the beads by shaking **5 min** at **room temperature**. Alternatively, resuspend beads completely by repeated pipetting up and down and incubate for **10 min** at **56** °C.

Separate the magnetic beads by placing the Square-well Block on the NucleoMag[®] SEP magnetic separator. Wait at least 2 min until all the beads have been attracted to the magnets. Transfer supernatant containing the purified genomic DNA to either microtubes or Tube Strips (see ordering information, section 6.2).

<u>Note:</u> Yield can be increased by 15-20 % by using prewarmed elution buffer (55 °C) or by incubating the bead / elution buffer suspension at 55 °C for 10 min.

6 Appendix

6.1 Troubleshooting

Problem	Possible cause and suggestions
	Elution buffer volume insufficient
	Beads pellet must be covered completely with elution buffer
	Insufficient performance of elution buffer during elution step
	 Remove residual buffers during the separation steps completely. Remaining buffers decrease efficiency of following wash steps and elution step.
	Beads dried out
Poor DNA yield	 Do not let the beads dry as this might result in lower elution efficiencies.
	Aspiration of attracted bead pellet
	 Do not disturb the attracted beads while aspirating the supernatant, especially when the magnetic pellet is not visible in the lysate.
	Aspiration and loss of beads
	• Prolong magnetic separation time or decrease aspiration speed.
	Insufficient washing procedure
Low purity	 Use only the appropriate combinations of separator and plate, for example, Square-well Block in combination with NucleoMag[®] SEP.
washing procedure. If shaking is not suf	 Make sure that beads are resuspended completely during the washing procedure. If shaking is not sufficient to resuspend the beads completely mix by repeated pipetting up and down.
	Carry-over of ethanol wash solutions
Suboptimal performance of	 Be sure to remove all of the 80 % ethanolic wash solution from the final wash, as residual ethanol interferes with downstream applications.
DNA in downstream	Ethanol evaporation from wash buffers
αρρισατοί ιδ	 Close buffer bottles tightly, avoid ethanol evaporation from buffer bottles as well as from buffer filled in reservoirs. Do not reuse buffers from buffer reservoirs.

	Time for magnetic separation too short
Carry-over of beads	 Increase separation time to allow the beads to be completely attracted to the magnetic pins before aspirating any liquid from the well.
-	Aspiration speed too high (elution step)
	 High aspiration speed during the elution step may cause bead carry-over. Reduce aspiration speed for elution step.
	Incomplete lysis
No or poor DNA yield	 Sample has not completely been submerged during heat incubation. Cut samples into small pieces. Mix well. Be sure that the samples are fully submerged in Buffer FL/Proteinase K mixture. Incubate until the samples are completely lysed.
	Reagents not applied properly
	• Prepare Buffer Proteinase K solution according to instructions (see section 3).
	RNA in sample
RNA contamination	 If DNA free of RNA is desired, cool down to room temperature after lysis incubation and add 20 µL of an RNase A solution (20 mg/mL; see ordering information, section 6.2). Incubate for 15 min with moderate shaking.
	Paraffin amount too high
Insufficient Paraffin removal	• Using to many paraffin sections or excess of paraffin may cause immediate solidification of the paraffin after heat incubation when cooling lysate down to room temperature. Repeat heat incubation with increased (e.g., doubled) volume of paraffin dissolver.

6.2 Ordering information

Product	REF	Pack of
NucleoMag [®] DNA FFPE	744320.1 744320.4	1 × 96 preps 4 × 96 preps
NucleoSpin [®] 96 DNA FFPE	740240.1 740240.4	1 × 96 preps 4 × 96 preps
NucleoSpin [®] FFPE DNA XS	740980.10 740980.50 740980.250	10 preps 50 preps 250 preps
Paraffin Dissolver (blue)	740343.60	60 mL
Proteinase K (lyophilized)	740506	100 mg
RNase A (lyophilized)	740505.50 740505	50 mg 100 mg
NucleoMag [®] SEP	744900	1
Square-well Blocks	740481 740481.24	4 24
Self adhering PE Foil	740676	50 sheets
Rack of Tube Strips (set consists of 1 Rack, 12 Tube Strips with 8 tubes each, and 12 CapStrips)	740477 740477.24	4 sets 24 sets
Elution Plate U-bottom	740486.24	24
Elution Plate Flat-bottom	740673	20
KingFisher [®] 96 Accessory Kit A (set consists of Square-well Blocks, Deep-well Tip combs, Elution Plates; for 4 × 96 NucleoMag [®] DNA FFPE preps using KingFisher [®] 96 platform)	744950	1 set

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

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.

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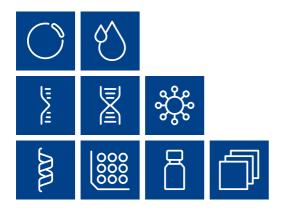
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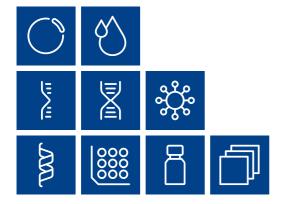
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Plasmid DNA Clean up RNA DNA Viral RNA and DNA Protein High throughput Accessories Auxiliary tools



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