

Instructions for use Invisorb® Fragment CleanUp

INVITEK
diagnostics





InviSorb®

Language: EN

RUO

REF 1020300300

 250 preparations

 ALS Life Sciences Portugal, S.A.
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Important notes

Thank you for purchasing the **InviSorb® Fragment CleanUp** from Invitek Diagnostics.

The **InviSorb® Fragment CleanUp** provides a convenient tool for fast and efficient purification of PCR products and DNA fragments from amplification or enzymatic reactions. Furthermore, this kit is the ideal tool for extraction of DNA fragments of 80 bp – 30 kb from standard or low melting agarose gels in TAE and TBE buffers at high final DNA concentrations. Up to 300 mg agarose gel slices can be processed per spin column.

WARNING! Improper handling and use for other than the intended purpose can cause danger and damage. Therefore, we ask you to read through these instructions for use and follow them carefully. Always keep them handy. To avoid personal injury, also observe the safety instructions.

All versions of the instructions for use can be found on our website for download or can be requested from us: www.invitek.com

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Kit contents of InviSorb® Fragment CleanUp

	250 preps
Catalogue No.	1020300300
Gel Solubilizer S	2 x 140 ml
Binding Buffer	63 ml (final volume 163 ml)
Binding Enhancer	30 ml (final volume 150 ml)
Wash Buffer	2 x 45 ml (final volume 2 x 150) ml)
Elution Buffer	15 ml
Spin Filter	5 x 50
2.0 ml Receiver Tubes	5 x 50
1.5 ml Receiver Tubes	5 x 50
Manual	1
Initial steps	<p>add 105 ml 96-100% ethanol to each bottle Wash Buffer</p> <p>add 120 ml 99.7% Isopropanol to the Binding Enhancer; Mix by intensive shaking for 1 min. Shortly before use mix by inverting several times</p> <p>add 100 ml 99.7% Isopropanol to the Binding Buffer; Mix by intensive shaking for 1 min. Shortly before use mix by inverting several times</p>

Symbols used on product and labeling

-  Manufacturer
-  Batch number
-  Catalogue number
-  Expiry date
-  See instruction manual
-  Temperature limitation
-  Do not reuse
-  Number of sample preparations
-  Research Use Only

Storage

All buffers and kit contents of the **InviSorb® Fragment CleanUp** should be stored at room temperature and are stable for at least 12 months.

Room temperature (RT) is defined as range of 15-30°C.

Before every use make sure that all components have room temperature. If there are any precipitates in the provided solutions, they can be dissolved by careful warming (up to 30°C).

Warranty

Invitek Diagnostics guarantees the correct function of the kit for applications described in this manual and in accordance with the intended use. In accordance with Invitek Diagnostics's EN ISO 13485 and ISO 9001 certified Quality Management System the performance of all kit components has been tested to ensure product quality.

Any problems, incidents or defects shall be reported to Invitek Diagnostics immediately upon detection. Immediately upon receipt, inspect the product to ensure that it is complete and intact. In the event of any discrepancies, you must inform Invitek Diagnostics immediately in writing. Modifications of the kit and protocols and use that deviate from the intended purpose are not covered by any warranty.

Invitek Diagnostics reserves the right to change, alter, or modify any product to enhance its performance and design at any time. Invitek Diagnostics warrants products as set forth in the General Terms and Conditions available at www.invitek.com. If you have any questions, please contact techsupport@invitek.com

Intended use

Before using the kit, please ensure that you have read the instructions and are fully informed about the purpose and limitations of use of the kit, and "Features of the InviSorb® Fragment CleanUp").

THE PRODUCT IS INTENDED FOR USE BY PROFESSIONALS ONLY, SUCH AS TECHNICIANS, PHYSICIANS AND BIOLOGISTS TRAINED IN MOLECULAR BIOLOGICAL TECHNIQUES. It is designed to be used with any downstream application employing enzymatic amplification or other enzymatic modifications of DNA followed by signal detection or amplification.

Product use limitation

For purification DNA fragments should not be bigger than 30 kb and not smaller than 80 bp. The maximum length of primers which can be removed is 40 bp. The InviSorb® Fragment CleanUp is not suitable for extraction of circular plasmids as these can be detected in multiple positions in the gel due to their very different configurations.

When carrying out the PCR with Taq DNA Polymerase, it is possible that the "A-overhangs" may be lost during extraction. It is therefore recommended to repair the A-overhangs if they are needed in the subsequent downstream application.

Safety instructions

Ensure that anyone using this product has received instructions in general safety practices for laboratories and the safety information provided in this document.

- When and while working with chemicals, always wear protective clothing, disposable gloves and safety glasses.
- Always change pipette tips between liquid transfers. To avoid cross-contamination, we recommend the use of aerosol-barrier pipette tips.
- Do not reuse any consumables.
- Discard gloves if they become contaminated.
- Do not combine components of different kits.
- Avoid microbial contamination of the kit reagents.
- To minimize the risk of infections from potentially infectious material, we recommend working under laminar airflow until the samples are lysed.

Before handling chemicals read and understand all applicable safety data sheets (MSDS). These are available online at www.invitek.com.

Dispose of kit residues and waste fluids in accordance with your country's regulations, again refer to the MSDS. Invitek Diagnostics has not tested the liquid waste generated by the kit for residual infectious materials. Contamination of the liquid waste with residual infectious materials is highly unlikely but cannot be excluded completely. Therefore, liquid waste must be considered infectious and must be handled and disposed of according to local safety regulations.

European Community risk and safety phrases for the components of the **InviSorb® Fragment CleanUp** to which they apply are listed below as follows:

Gel Solubilizer S



Danger

Contains : guanidinium thiocyanate

Hazard statements

H302+H312 - Harmful if swallowed or in contact with skin.

H314 - Causes severe skin burns and eye damage.

H412 - Harmful to aquatic life with long lasting effects

Precautionary statements

P260 - Do not breathe dust/fume/gas/mist/vapours/spray.

P273 - Avoid release to the environment.

P301+P312 - IF SWALLOWED: Call a POISON CENTRE or doctor if you feel unwell.

P301+P330+P331 - IF SWALLOWED: Rinse mouth. Do NOT induce vomiting.

P302+P352 - IF ON SKIN: Wash with plenty of water.

P303+P361+P353 - IF ON SKIN (or hair): Take off immediately all contaminated clothing. Rinse skin with water.

P501 - Dispose of contents/container to hazardous or special waste collection point, in accordance with local, regional, national and/or international regulation
P260 - Do not breathe dust/fume/gas/mist/vapours/spray.

Emergency medical information can be obtained 24 hours a day from infotrac:

outside of USA: 1 – 352 – 323 – 3500

in USA: 1 – 800 – 535 – 5053

PCR product & DNA fragment purification and concentration

This manual describes one kit for both, the membrane adsorption-based purification via high-performance MSB[®] technology of PCR products and products from other enzymatic reactions as well as the purification of DNA fragments from agarose gels using the InviSorb[®] spin columns via the standard lyse-bind-wash-elute-protocol.

	Gel Solubilization Technology
Sample Volume	up to 100 µl * or up to 300 mg gel slices
Recovery	80 – 95 %* 60 – 90 %
Binding capacity	10 µg
Elution Volume (minimal)	10 µl */ 20 µl
Sample Source :	
- PCR reaction mixture	x*
- Ligation reaction mixture	x*
- Enzyme digestion mixture	x*
- cDNA synthesis mixture	x*
- Cycle sequencing reaction	x*
- DNA fragments	x*
- Agarose gels (TAE, TBE)	x

* MSB[®] technology

Advantages

- convenient and fast sample processing
- the most efficient removal of contaminants
- high recovery rate of PCR products or DNA fragments
- broad range of fragment sizes: 80 bp - 30 kb can be purified

The spin columns are designed to give high final concentrations of purified DNA fragments for subsequent reactions. Special buffers provided with this kit are optimized for efficient recovery of DNA and removal of contaminants like, salts, enzymes, nucleotides, agarose, ethidium bromide and other impurities from DNA samples. Specialized binding buffers promote selective adsorption of DNA fragments and PCR products. The pure DNA is eluted in a small volume of buffer or water, ready to use for any subsequent application. The innovative **MSB® technology** is described in the following chapter.

MSB® technology

The fastest technology for purification of DNA fragments with high recovery rates.

The MSB® technology offers a new option for the purification process in which handling steps are greatly simplified and processing times are significantly reduced. The products with **MSB® technology** have been designed for extremely efficient purification and/or concentration of PCR products and DNA fragments from enzymatic reaction mixtures in only two steps.

The DNA fragments adsorb at the silica membrane in the presence of minimal concentrations of non-chaotropic salts, while impurities pass through the column. Therefore, a washing step is not required. High concentrated, pure DNA fragments are eluted ready for use.

Advantages:

- ultra-fast and easy (two step protocol), only binding and elution
- excellent purity without washing
- 80 – 95 % recovery rate

DNA purified with the MSB® technology is much more concentrated than DNA purified by other methods. The highly concentrated DNA allows the use of small reaction volumes, which are useful for any downstream application, leading to increased efficiency (e.g. in ligations).

Features of the InviSorb® Fragment CleanUp

Starting material Size of DNA fragments: 80 bp - 30 kb	Yield	Time for preparation
up to 100 µl reaction volume like PCR reaction mixture, up to 100 µl restriction digestion mixture, up to 100 µl ligation mixture, up to 100 µl cDNA synthesis mixture, up to 100 µl cycle sequencing reaction mixture	80 – 95 %, depends on fragment length	approx. 7 min
up to 300 mg of gel slices (0.8 – 2%) from TAE or TBE agarose gels	60 - 90 %, depends on fragment length and kind of agarose gel	approx. 20 min

The **InviSorb® Fragment CleanUp** provides ultra-fast purification and concentration of up to 100 µl PCR products, or of linearized plasmid DNA as well as purification of DNA fragments from agarose gels and clean-up of other enzymatic reactions.

For PCR purification the MSB® method is used, which requires no washing steps, DNA fragments are eluted with low salt buffer or water.

For purification of DNA-fragments from agarose gels, the standard lyse-bind-wash-elute-protocol is used. DNA fragments bind directly to the surface of a spin filter column after gel solubilization. After washing steps, elution of the DNA fragments is performed with a low salt buffer.

While the MSB® method works without chaotropic salts, chaotropic salts are necessary for lysis from an agarose gel. Therefore, depending on the protocol, both types of chemistry, chaotropic or non-chaotropic, are used for DNA binding in this kit.

The purified DNA-fragments are ready to use in various downstream applications such as:

- digestion with restriction enzymes
- hybridization, labelling, cloning
- sequencing
- *In vitro* transcription
- ligation and transformation
- DNA sequencing
- amplification

Before starting a protocol

Check the product and its components as well as the packaging for obvious damage, correct quantities and quality immediately upon receipt of the product.

- Always change pipette tips between liquid transfers. To avoid cross-contamination the use of filter tips is recommend.
- All centrifugation steps are carried out at room temperature.
- When working with chemicals, always wear a lab coat, disposable gloves and protective goggles.
- Discard gloves if they become contaminated.
- Do not mix kit components with components from other kits unless the lot numbers are identical.
- Avoid microbial contamination of the kit reagents.
- To minimize the risk of infections from potentially infectious material, we recommend working under laminar air-flow until the samples are lysed.

Equipment and reagents to be provided by the user

- Microcentrifuge ($\geq 11.100 \times g$)
- Ethanol (96-100%)
- Thermoshaker
- Pipettes and filter tips
- Scalpel 120
- 1.5 ml and 2.0 ml reaction tubes
- Isopropanol (99.7%)

Principle and procedure of the InviSorb® Fragment CleanUp

The InviSorb® Fragment CleanUp combines two applications: DNA fragment purification and agarose gel extraction.

Procedure for DNA fragment purification:

- selective binding of DNA fragments to the surface of the DNA-Binding Spin Filter
- elimination of enzyme buffer, enzyme, primers and nucleotides during the binding step
- elution of the highly pure DNA fragment

Sampling and storage of starting material

Best results are obtained using freshly prepared PCR or enzymatic reaction mixtures to prevent DNA digestion. The samples can be stored for some weeks at 4 - 8°C.

Binding of DNA fragments

The reaction mixture is mixed in a ratio of 1:5 with the **Binding Buffer** to provide the appropriate condition for the binding of DNA fragments in a range of 80 bp - 30 kb to the silica membrane under minimal concentrations of non-chaotropic salts.

The binding of small DNA fragments can be supported by the addition of small amounts of isopropanol, but this ratio is very sensitive.

Removal of Contaminants

The DNA fragments bind to the membrane at minimal concentrations of non-chaotropic salts. Therefore, a washing step is not required. Unwanted primers and impurities such as salts, enzymes, unincorporated nucleotides, dyes, ethidium bromide, oils, and detergents do not bind to the silica membrane; instead, they are pulled through the column by centrifugal force together with the large excess of Binding Buffer. Any remaining Binding Buffer, which may interfere with subsequent enzymatic reactions, is removed by additional centrifugation.

Elution of PCR products or DNA fragments

DNA is eluted from the column using 10 - 50 µl Elution Buffer.

Eluting twice each with 30 - 50 µl leads to complete recovery of DNA. By the use of smaller elution volumes DNA concentration can be increased. Elution volumes should not fall below 10 µl, otherwise the yield will be reduced. The eluted DNA is ready to use in different downstream applications.

Procedure for gel extraction:

- excision of the DNA-fragment from the agarose gel with a sharp scalpel
- gel removal and binding of DNA fragments on the membrane of the spin column
- selective binding of DNA fragments to the surface of the DNA-Binding Spin Filter
- elimination of contaminants and ethanol
- elution of the highly pure DNA fragment

Sampling and storage of starting material

Use low melting and standard gels (0.8 – 2%) with TAE or TBE buffer systems. Best results are obtained using freshly prepared DNA slices from a gel. However, the samples can be stored for some weeks at 4 - 8°C.

Excision of DNA fragments

For best results, the pieces of gel should be as small as possible. Do not expose the gel with the DNA fragment to UV light for a long time, reduce cutting time under UV light to a minimum since it is damaging to the DNA**.

Gel removal and binding of DNA fragments

Gel Solubilizer S in the **InviSorb® Fragment CleanUp** solubilizes the agarose gel slice under high temperatures. Together with Binding Enhancer and Binding Buffer, it provides the appropriate condition for the binding of the DNA to the silica membrane at high salt concentrations.

Removal of contaminants and of Ethanol

The DNA fragments bind to the membrane, contaminants and salts are washed away by the Ethanol-containing Wash Buffer. Any remaining Wash Buffer, which may interfere with subsequent enzymatic reactions, is removed by additional centrifugation.

Elution of PCR products or DNA fragments

DNA is eluted from the column using 10 - 50 µl Elution Buffer.

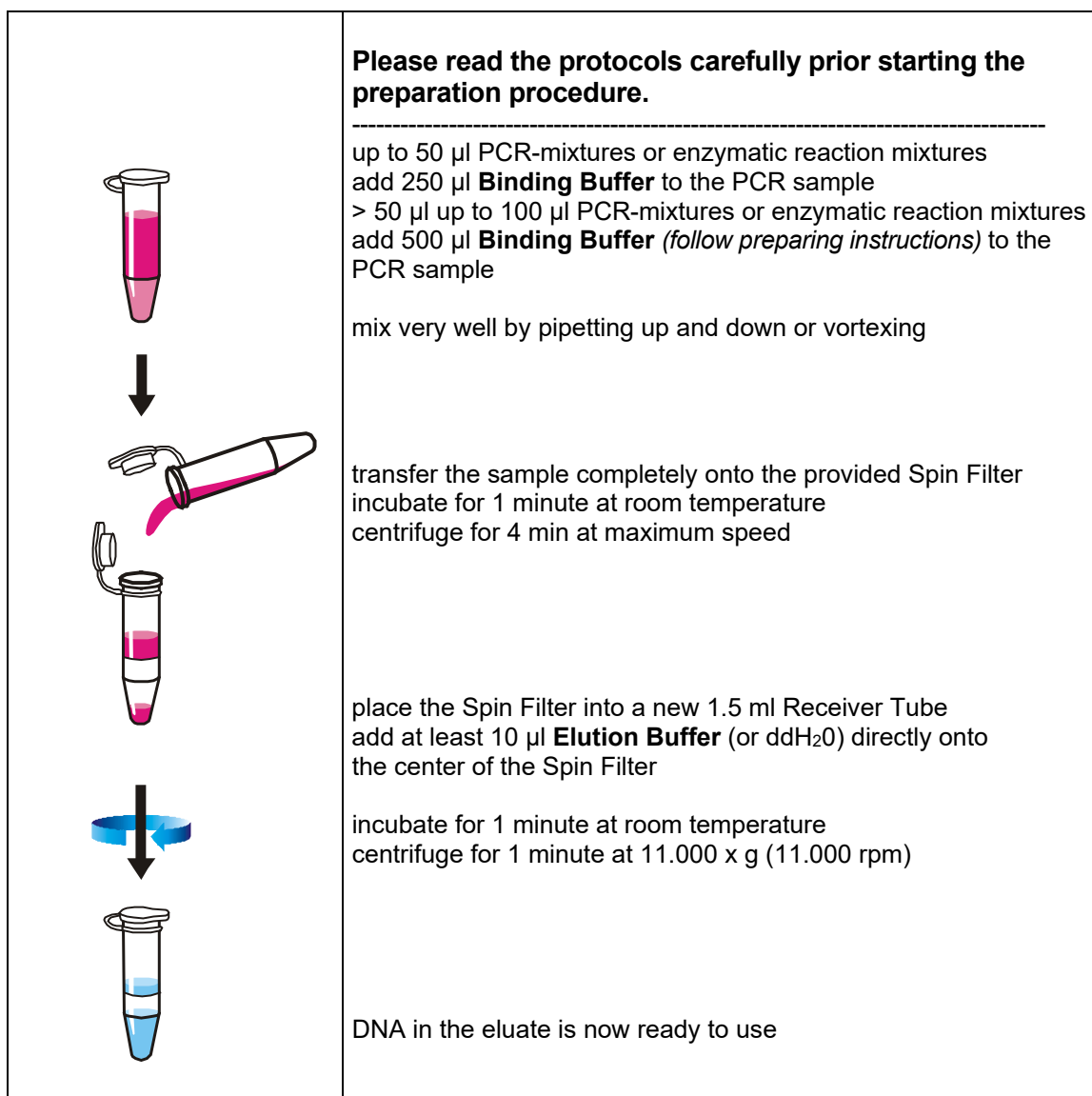
Eluting twice each with 30 - 50 µl leads to complete recovery of DNA. By the use of smaller elution volumes DNA concentration can be increased. Elution volumes should not fall below 10 µl, otherwise the yield will be reduced. The eluted DNA is ready to use in different downstream applications.

**) see protocol 5

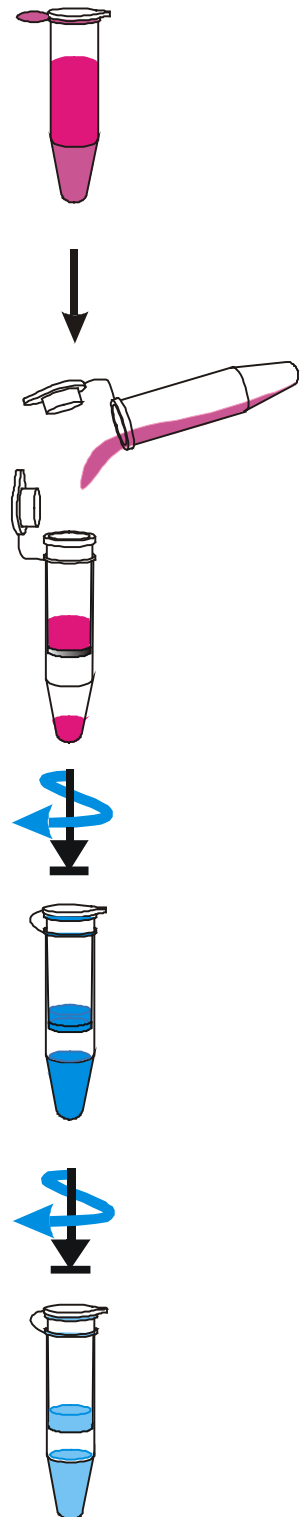
Preparing reagents and buffers of the InviSorb® Fragment CleanUp

250 preps
Add 105 ml 96-100% Ethanol to each bottle Wash Buffer Add 120 ml 99.7% Isopropanol to the Binding Enhancer ; Mix by intensive shaking for 1 min. Shortly before use mix by inverting several times Add 100 ml 99.7% Isopropanol to the Binding Buffer ; Mix by intensive shaking for 1 min. Shortly before use mix by inverting several times

Scheme for DNA fragment purification



Scheme for agarose gel extraction

	<p>Please read the protocols carefully prior starting the preparation procedure.</p> <hr/> <p>Transfer gel slices (max. 300 mg) into a 1.5 ml or 2.0 ml microcentrifuge tube (not provided)</p> <p>to gel slices up to 150 mg add 500 μl Gel Solubilizer S to gel slices > 150 mg add 1 ml of Gel Solubilizer S</p> <p>incubate at 50°C for 10 minutes until the gel is completely solubilized</p> <p>add 250 μl Binding Enhancer (<i>follow preparing instructions</i>) to a 500 μl reaction volume add 500 μl Binding Enhancer (<i>follow preparing instructions</i>) to a 1 ml reaction volume</p> <p>mix by pipetting up and down for 2-3 times</p> <p>load approx. 800 μl of the sample onto the Spin Filter centrifuge at 11.000 x g (11.000 rpm) for 2 min discard the filtrate for reaction volumes > 800 μl reload the residual volume and repeat the centrifugation step</p> <p>add 500 μl Wash Buffer to the Spin Filter centrifuge for 1 min at 11.000 x g (11.000 rpm) discard the filtrate repeat the washing step once again centrifuge for 1 min at 11.000 x g (11.000 rpm), discard the filtrate</p> <p>remove the remaining ethanol by centrifugation for 4 min at maximum speed</p> <p>transfer the Spin Filter into a new 1.5 ml Receiver Tube add at least 10 μl Elution Buffer directly to the center of the Spin Filter incubate at room temperature for 1 min centrifuge for 1 minute at 11.000 x g (11.000 rpm)</p> <p>DNA is now ready to use</p>
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Protocol 1: Purification and concentration of DNA fragments from enzymatic reactions, e.g. PCR-products from PCR reactions, cDNA synthesis, enzyme restriction digestions

Please read the instructions carefully and carry out preparatory arrangements in advance.

Note: Before starting with the purification procedure please place a Spin Filter into a 2.0 ml Receiver Tube!

Attention: Please prepare the **Binding Buffer** ahead- see instruction page: 11

1. Binding of the PCR or DNA - fragments

A. For PCR-mixtures up to 50 µl

Add **250 µl Binding Buffer** to the PCR sample and mix very well by pipetting or vortexing. Transfer the sample completely to a Spin Filter, incubate for 1 minute at room temperature and centrifuge for 4 min at maximum speed.

B. For PCR-mixture > 50 µl up to 100 µl

Add **500 µl Binding Buffer** to the PCR sample and mix very well by pipetting or vortexing. Transfer the sample completely onto a Spin Filter, incubate for 1 minute at room temperature and centrifuge for 4 min at maximum speed.

2. Elution of the PCR or DNA - fragments

Place the Spin Filter into a new 1.5 ml Receiver Tube.
Add at least 10 µl Elution Buffer (or ddH₂O) directly onto the center of the Spin Filter.
Incubate for 1 minute at room temperature. Centrifuge for 1 minute at 11.000 x g (11.000 rpm).

Important notes:

- 1. If the PCR-mixture contains mineral oil, we recommend the addition of 500 µl of Binding Buffer independent of the starting volume. It is also possible to wash the bound PCR-fragment once with 500 µl of Binding Buffer.*
- 2. To increase the final DNA yield we recommend an extended incubation time with Elution Buffer (up to 5 minutes), which will lead to a slightly higher final yield.*
- 3. For concentration of PCR-fragments it is possible to elute with lower volume of Elution Buffer, than the volume of the starting PCR-mixture. The minimum volume is 10 µl.*
- 4. For ligation mixtures please note, that ligation reactions give very often unwanted side products. These are purified and enriched as well.*

Protocol 2: Removal of DyeDeoxy™ terminators from DNA cycle sequencing reactions of PCR-products and plasmids after use ABI Prism™ terminator Kits

Please read the instructions carefully and carry out preparatory arrangements in advance.

Note: Before starting with the purification procedure please place a Spin Filter into a 2.0 ml Receiver Tube!

Attention: Please prepare the **Binding Buffer** ahead- see instruction page: 11

1. Binding of the (fluorescent) labeled DNA

Add **500 µl Binding Buffer** to the completed cycle sequencing reaction (up to 100 µl) and mix thoroughly by pipetting or vortexing. Transfer the sample completely onto a Spin Filter and centrifuge for 4 min at maximum speed.

Note:

If sequences next to the primer (short fragments) shall be obtained, the addition of up to 150 µl of Isopropanol to the upper mixture may be helpful (the shorter the desired fragments are the more Isopropanol must be used). This leads to lower purity but also to recovery of shorter fragments.

2. Elution of the (fluorescent) labeled DNA

Place the Spin Filter into a new 1.5 ml Receiver Tube.

Add at least 10 µl Elution Buffer (or ddH₂O) directly onto the center of the Spin Filter.

Incubate for 1 minute at room temperature. Centrifuge for 1 minute at 11.000 x g (11.000 rpm).

Discard the Spin Filter and proceed with the ABI sample loading.

Protocol 3: Extraction of DNA fragments from agarose gel slices

Please read the instructions carefully and carry out preparatory arrangements in advance.

Important: *TBE-gels contain more potentially inhibitors for downstream application than TAE-gels. Therefore, we recommend the use of TAE-gels for critical downstream applications! Before starting with the purification procedure please place a Spin Filter into a 2.0 ml Receiver Tube!*

Attention: *Please prepare the **Binding Buffer** ahead - see instruction page: 11*

1. Excise the DNA-fragment from 0.8 – 2% agarose gel with a sharp scalpel. The gel piece should be cut out as small as possible. Prevent long exposure to UV light. Verify the weight of the piece.

**For gel slices up to 150 mg add 500 µl Gel Solubilizer S.
For gel slices > 150 mg – 300 mg add 1 ml of Gel Solubilizer S.**

Do not use more than 300 mg gel slice for one Spin Filter.
Transfer the gel slice into a 1.5 or 2.0 reaction tube.

2. Incubate at 50°C for 10 minutes until the agarose gel slice is completely dissolved. Continuous shaking during incubation (e.g. Eppendorf Thermo mixer) is very helpful.
3. Add 250 µl Binding Enhancer to a 500 µl reaction volume or 500 µl Binding Enhancer to a 1 ml reaction volume and mix the suspension by pipetting or by vortexing. Load approx. 800 µl of the sample onto the Spin Filter. Centrifuge at 11.000 x g (11.000 rpm) for 2 minutes. Discard the filtrate. For reaction volumes > 800 µl reload the remaining volume onto the Spin Filter and repeat the centrifugation step.
4. Add 500 µl Wash Buffer to the Spin Filter and centrifuge for 1 min at 11.000 x g (11.000 rpm). Discard the filtrate. Repeat the washing step once again.
5. Discard the filtrate. Remove the residual ethanol of the Wash Buffer by centrifugation for 4 min at maximum speed.
6. Transfer the Spin Filter to a new 1.5 ml Receiver Tube. Add at least 20 µl Elution Buffer directly to the center of the Spin Filter. Incubate at room temperature for 1 minute. Centrifuge for 1 minute at 11.000 x g (11.000 rpm).

Note: *To increase the final DNA yield we recommend using a higher volume of Elution Buffer. Please take into account that an increasing volume of Elution Buffer reduces the final concentration of the purified DNA. An extended incubation time with Elution Buffer (up to 10 minutes) leads also to a slightly higher final yield.*

Additional Protocol 4: Purification of PCR - products from 200 µl PCR reactions

Please read the instructions carefully and carry out preparatory arrangements in advance.

Note: Before starting with the purification procedure please place a Spin Filter into a 2.0 ml Receiver Tube!

Attention: Please prepare the **Binding Buffer** ahead- see instruction page: 11

1. Binding of the PCR-fragments

For PCR-reaction of 200 µl

Add **1000 µl Binding Buffer** to the PCR sample and mix thoroughly by pipetting or vortexing. Transfer the sample in two aliquots onto a Spin Filter and centrifuge for 1 min at 11.000 x g (11.000 rpm) each. Remove the filtrate and centrifuge again for 4 minutes at maximum speed.

2. Elution of the PCR-fragments

Place the Spin Filter into a new 1.5 ml Receiver Tube.

Add at least 10 µl Elution Buffer (or ddH₂O) directly onto the center of the Spin Filter.

Incubate for 1 minute at room temperature. Centrifuge for 1 minute at 11.000 x g (11.000 rpm).

Important Notes:

- 1. The provided volume of Binding Buffer is calculated based on the required buffer volumes in protocol 1 and 2. The amount needed for protocol 4 is not considered.*
- 2. If the PCR-mixture contains mineral oil, we recommend the addition of 500 µl Binding Buffer independent of the starting volume. It is also possible to wash the bound PCR fragment once with 500 µl of Binding Buffer.*
- 3. To increase the final DNA yield we recommend an extended incubation time with Elution Buffer (up to 5 minutes), which will lead to a slightly higher final yield.*
- 4. For concentration of PCR-fragments, it is possible to elute with lower volume of Elution Buffer than the volume of the starting PCR-mixture. The minimum volume is 10 µl.*

Additional Protocol 5: Instruction for adding A-overhangs to PCR products after gel purification

Please read the instructions carefully and carry out preparatory arrangements in advance.

Transfer 30 µl of the extracted PCR product to a 1.5 ml reaction tube.

Add 2 units standard Taq DNA Polymerase (no proofreading activity), 3.5 µl Taq Buffer, 0.6 µl dNTPs (10 mM each), if not contained in the Taq Buffer in a 35 µl reaction volume

Incubate for 15 min at 72°C under continuous shaking in a thermomixer.

After this treatment fragments can be used in cloning/ligation experiments.

Troubleshooting for agarose gel extraction

Problem	Cause	Comments and suggestions
Low recovery	No ethanol added to the Wash Buffer	Prepare the Wash Buffer exactly as described in the manual. Store the Wash Buffer with firmly fixed lid.
	Poor elution of DNA, ineffective solubilization of the agarose gel slice	Add the Elution Buffer directly onto the center of the Spin Filter (also if a small elution volume is used). For smaller fragments than 500 bp, please use TAE agarose gels.
Problems with down stream application, e.g. ligation	No Binding Enhancer added	Avoid this mistake.
	Contamination with salt components Contamination with agarose traces	The gel slice must be completely dissolved. Add the amount of Binding Enhancer needed to the solubilized suspension. Wash off the Spin Filters as described in the manual. Prolong the incubation time with Wash Buffer to 5 minutes before centrifugation.
	Contamination of the final DNA with ethanol	Keep the given centrifugation time, extend it if necessary (verify the smell).

Troubleshooting for DNA fragment purification

Problem	Cause	Comments and suggestions
Low recovery	Poor elution of DNA	Add the Elution Buffer directly onto the center of the Spin Filter (even if a small elution volume is used).
	Problems with mineral oil	Try to avoid pipetting of mineral oil. Apply the correct centrifugation steps. Take a higher volume of Binding Buffer . Wash once with Binding Buffer .

Ordering information

Product	Package Size	Catalogue No.
InviSorb® Fragment CleanUp	250 purifications	1020300300

Revision history

Revision	Date	Description
DE 573.01	2025-07-31	New document



INVITEK diagnostics

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