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### Chapter 11 Index

#### Chromatography

Chromatography methods may utilize gel filtration, ion exchange, selective adsorption, or affinity binding.

*Gel filtration* exploits the molecular sieving properties of porous gel particles. A matrix with defined pore size allows smaller molecules to enter the pores by diffusion whereas bigger molecules are excluded from the pores and eluted at the void volume. Thus, molecules are eluted in order of decreasing molecular size.

*Ion exchange chromatography* depends on an electrostatic interaction between a target molecule and a functional group on the column matrix. The technique allows concentration and separation of molecules from a large volume in a short time. Nucleic acids – highly negatively charged, linear polyanions – can be eluted from ion exchange columns with simple salt buffers.

In *adsorption chromatography*, nucleic acids adsorb selectively onto silica or glass in the presence of chaotropic salts, while other biological molecules do not. A low salt buffer or water then elutes the nucleic acids, thereby producing a sample that could be used directly in most downstream applications.

*Affinity chromatography* is a highly specific adaptation of adsorption chromatography. An immobilized ligand recognizes and binds a particular structure on a biomolecule. Washes then remove unbound components (with different structures). Finally, a "competitor molecule" (which also recognizes the immobilized ligand) floods the binding sites on the affinity matrix, releasing the bound biomolecule.

#### Centrifugation

Selective centrifugation is a powerful purification method. For example, u tr centrifugation in self-forming CsCl gradients at high g-forces has long by Cased for plasmid purification.

Frequently, centrifugation is combine with a lotter method. I on example:

• Spin column chromatography combines gel fileration and centrifugation to purify DNA or RNL from smaller contaminants (al.s, nucleotides, etc.), for buffer ex-

Some proced was of the selective adsorption on a chromatographic matrix (see above) with centrifugal elution to selectively purify one type of nucleic acid.

#### Electrophoresis

Nucleic acids may be separated electrophoretically according to their size. This separation is most commonly done on agarose gels. In the presence of ethidium bromide, the separated nucleic acids may be seen under UV light.

Electrophoretic separation is also frequently used to determine size and purity of DNA. For example, after PCR, electrophoresis is used to quickly check product length and purity (absence of byproducts).

#### Affinity purification

In recent years, more and more purification methods have combined affinity immobilization of nucleic acids with magnetic separation. For instance,  $poly(A)^+$  mRNA may be bound to streptavidin-coated magnetic particles by biotin-labeled oligo(dT) and the particle complex removed from the solution (and unbound contaminants) with a magnet. This solid phase technique simplifies nucleic acid purification, since it can replace several centrifugation, organic extraction, and phase separation steps with a single, rapid magnetic separation step.

### Principle of silica adsorption

All the kits described in this chapter depend on the tendency of nucleic acids to adsorb to silica (glass) in the presence of a chaotropic salt such as sodium iodide (NaI), guanidine thiocyanate or guanidine hydrochloride (Melzak et al., 1996).

This tendency was discovered by Vogelstein and Gillespie (1979) who found that DNA fragments adsorbed to powdered flint glass in the presence of saturated NaI. Later work showed other nucleic acids adsorbed to glass in the presence of other chaotropes, including:

- DNA plasmids (Marko et al., 1982)
- Single-stranded phage nucleic acids (Kristensen et al., 1987; Zimmermann et al., 1989)
- Genomic DNA (Yamada et al., 1990; Zeillenger et al., 1993)
- Total RNA (Yamada et al., 1990)
- Rapid and simple method for purification of nucleic acids (Boom et al., 1990)

Different types of nucleic acid adsorb more or less tightly to glass depending on the ionic strength and the pH of the surrounding solution. A low salt buffer or water is always used to elute the nucleic acid from the glass. In each kit, this method is optimized to prepare a particular type of nucleic acid.

#### **Overview of the procedure**

,co.u ecta plastic Filter Tube. All the High Pure kits use glass fiber fleece immobilized in The glass fleece filter:

Adsorbs only nucleic acid lecules from a baration of the rin m xt in of proteins, sug ipid complex biolog other components

tube and processed in a standard tabletop ed into a microcentr

- Allows processing of 0.2 2.5 ml samples in a series of centrifugation steps
- Is specially constructed to ensure that contaminants suspended in the Wash Buffer are not retained by the filter or transferred to the eluted, purified nucleic acid

#### Benefit from smart column design

#### The High Pure Micro Column Achieve high purity. Novel optimized columns produce highly concentrated (10 µl) eluate and high recovery rates (>80%) of even small DNA and RNA fragments. • Up to 10 µg binding capacity for use in demanding downstream applications. Avoid carryover contamination using a column design without a cavity for liquid transfer. The High Pure Mini Column Perform long template applications. Efficiently purify high molecular weight DNA (30-50 kb). Maximize performance and accuracy in downstream assays. Achieve high sensitivity and reproducibility in many applications with a binding capacity of up to 100 µg. Obtain accurate results. Use highly pure, concentrated (50 µl) nucleic acids in real-time PCR and other applications.

#### The High Pure Extender Assembly

- Benefit from smart column design using a removable High Pure mini spin column.
- S.
- Improve sensitivity. Use sample volumes up to 2.5 ml for high yields of purified nucleic acids concentrated in 50 µl.
   Obtain high-purity nucleic acids. Reduce carryover risk by using high centrifugal
- orces in all wash steps.
   Increase convenience and improve time to result. Eliminate complicated
- sample pre-processing and rapidly recover purified nucleic acids using high-speed centrifugation.

In each High Pure kit, the steps are basically the same and require only a few minutes. The nucleic acids prepared with the High Pure kits may be used directly in a variety of down-stream applications.

Substance added to serum	Highest concentration tested with no inhibition in PCR after High Pure purification
Citrate	30 mg/ml
EDTA	300 mg/ml
Heparin	30 U/ml
Hemoglobin	50 mg/ml

Performance of High Pure Nucleic Acid Purification in the removal of different anticoagulants or human hemoglobin. The table indicates the highest concentration tested which showed no inhibiting.

Instead of glass fiber fleece (as in the High Pure kit), one capalso use silica beads to adsorb DNA.

The Agarose Gel DNA better icht at starts with an agarose gel slice containing a DNA fragment a el solubilizes that gel to de a set le DNA into the starting solution

The lit uses steps similar to the of the digh Pure kits to purify the DNA from the starting material. In each, the of cleic acid is adsorbed to silica in the presence of a chaotropic with the lited by centrifugation (while adsorbed to the silica beads), washed extensively or remove contaminants, then released from the beads with a low salt buffer.

The DNA isolated with the kit is pure enough to be used directly in labeling, sequencing, cloning, and other procedures that require concentrated DNA.

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Typical results with the kit

#### **Experiment 3**



Figure 3: Use of nucleic acids from paraffin-embedded tissue (prepared with the High Pure PCR Template Preparation Kit) for amplification of the human androgen receptor gene. Nucleic acids were purified from formalin-fixed, paraffin-embedded samples of human prostate tissue by Protocol Ve. An aliquot (10  $\mu$ ) of each nucleic acid product was used as template for the Expand High Fidelity PCR System (100  $\mu$ l reaction mixture). The primers were derived from exon 7 of the androgen receptor gene. Aliquots (20  $\mu$ l) of each amplicon were separated electrophoretically and the gel was stained with ethidium bromide.

Lane 1: Molecular Weight Marker VIII Lane 2: Androgen receptor amplicon from one tissue section

**Lane 3:** Androgen receptor amplicon from half a tissue section

**Result:** The 263 bp androgen receptor amplicon was clearly visible in both tissue samples.

	Experiment 4		-0	uk
			Incupa () n p v oa	
		15 min	30 min	60 min
	L. casei	NOU		
	Without Mu	5.2 (40.0 %)	<b>7</b> . <b>1</b> 3.2 %)	21.8 (57.4 %)
	0.25 J/µ Wu	153 (40.37)	24.6 (64.7 %)	24.0 (63.2 %)
	ρ Vµl Mu	6 - (43.2 %)	38.0 (100 %)	30.4 (80.0 %)
nrev	L. curvetus			
PIC -	Without Mr	9.2 (10.6 %)	14.7 (17.0 %)	26.1 (30.2 %)
	0.25 U7µl Mu	25.9 (29.9 %)	61.3 (70.9 %)	70.7 (81.7 %)
	0.5 U/µl Mu	46.0 (53.2 %)	86.5 (100 %)	131.1 (151.6 %)
	L. sakei			
	Without Mu	13.5 (11.4 %)	16.6 (14.0 %)	52.9 (44.7 %)
	0.25 U/µl Mu	35.0 (29.6 %)	71.8 (60.6 %)	168.2 (142.1 %)
	0.5 U/µl Mu	79.3 (67.0 %)	118.4 (100 %)	196.1 (165.6 %)
	S. aureus			
	Without Ly	10.3 (10.8 %)	30.5 (31.9 %)	64.2 (67.2 %)
	10 µg/ml Ly	16.5 (17.3 %)	95.0 (99.5 %)	105.2 (110.2 %)
	25 μg/ml Ly	16.2 (17.0 %)	95.5 (100 %)	106.7 (111.7 %)

Table 1: Use special lysis conditions for gram-positive bacteria to achieve high DNA concentrations ( $\mu$ g/ml) in eluate. The Lysis Buffer contained 10 mg/ml lysozyme for lactobacilli and 1 mg/ml lysozyme for staphylococci together with the amounts of mutanolysin (MU) or lysostaphin (Ly) shown in the first column. The percentages refer to the DNA yield obtained with the highest mutanolysin or lysostaphin concentration and an incubation period of 30 min.

2

### LightCycler<sup>®</sup> sample preparation

The High Pure PCR Template Preparation Kit has been evaluated for LightCycler® sample preparation with whole blood and cultured cells as sample material. Amplification has been performed in LightCycler® capillaries using SYBR Green and Hybridization Probes as the detection formats. The following tables give information about the range of sample volume applied, modifications in the sample preparation procedure and expected results.

Sample	Volume/Amount Range	Typical concentration range [ng/µl]	Eluate use in LightCycler® PCR (µl)
Whole blood human*	1 – 50 µl **	0.5 – 25	1 – 5
			[0.5 ng – 125 ng]
Cultured cells, K562*	100 – 10⁵ cells	0.01 – 10	1 – 5
			[0.01 – 50 ng]

Research samples

\*\* Yield may vary between different blood donors because they may have different amounts of leukocytes

### **Procedure modification**

The standard procedure for whole blood and cultured cells has been used for LightCycler® sample preparation except the elution volume was set to 50 u. In order to increase nucleic acid concentration for minute sample amounts Por larger sample amounts the standard elution volume of 200 u.c. no opplied.

		. 1	<u>162</u>			
Parameter Sample SYBR Green				HybProbes		
		ົ່ງໃ	<b>N</b> n	Max	Min	Max
- ovi	cyclophilin A	Ploce	0.005 μl <sup>a)</sup>	100 µl	0.005 µlª)	100 µl
PIC.	or $\beta$ -, le jin	cells	1 <sup>a)</sup>	104	1 <sup>a)</sup>	10 <sup>5</sup>
<b>—</b>						

The above table shows the minimal and maximal sample amounts which have been purified and used for LightCycler® Amplification with whole blood and cultured cells as sample material. The parameters human cyclophilin A and  $\beta$ -globin have been investigated. All values are for 20  $\mu$ l LightCycler® Amplification when 5  $\mu$ l of 50  $\mu$ l total eluate is applied.

a) The used quantities of eluate correspond to these calculated amounts.

Time required	The entire High Pure PCR Cleanup Micro Kit method takes approx. 10 min.
Results	The amount of DNA recovered is dependent on the amount of DNA applied to the glass fiber fleece, the elution volume, and the length of the amplification/DNA products. When 5 - 25 µg DNA is applied to the kit's High Pure Micro Filter Tube, approximately 80% of the DNA can be recovered.
Benefits	• Conserve resources by using one versatile kit that eliminates the need to use several kits from other suppliers.
	• Save time with a simple and rapid protocol that reduces purification time.
	• Obtain purified product in a small elution volume (≤10 µl) for demanding down- stream applications.
	• Efficiently remove contaminants and unwanted reaction components.
	• Generate contaminant-free DNA for direct use in cloning, ligation, restriction digests, and other reactions.
	• Selectively isolate specific DNA fragment sizes by using the kit's binding enhancer to adjust purification stringency.

2

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### How to use the kit





### II. Kit contents

- Binding Buffer 20 ml or 80 ml
- Binding Enhancer 15 ml or 45 ml
- Wash Buffer 10 ml or 2 x 20 ml
- Elution Buffer 40 ml
- High Pure Micro Filter Tubes (50 or 200 tubes)
- Collection Tubes (50 or 200 tubes)

.

Typical results with the kit

$\frown)$

Vc. Purifying labeled probes	

1	Fill up the labeling reaction to 100 $\mu$ l with PCR grade water.
2	Add 500 µl Binding Buffer and mix well.
	It is important that the volume ratio between sample and Binding Buffer is 1:5. When using other sample volumes than $100 \mu$ l, adjust the volume of Binding Buffer accordingly.
3	Follow Protocol Va starting at the sample transfer to the Filter Tube assembly (Step 2).
	The elution efficiency is increased with higher volume of Elution Buffer applied. At least 68% and 79% recovery are found with 50 and 100 μl Elution Buffer, respectively. Normally, almost quantitative recovery can be found, as can be

### VI. Troubleshooting the High Pure protocols

determined in a direct detection assay.

The same troubleshooting procedure can be applied to all High Pure kits. For details on how to troubleshoot the above protocols, see the General Troubleshooting Procedure for all High Pure kits on page 113 of this manual. For factors that may affect the High Pure PCR Product Purification Kit, see page 116.



### Figure 8: Comparison of PCR products purified with the High Pure PCR Product Purification Kit and those purified by phenol/chloroform extraction.

Four different long PCR products were generated with the Expand Long Template PCR System, purified by either of two methods, then analyzed electrophoretically. From left to right, the four products shown on the gel are: a 2.9 kb fragment from the p53 gene; 4.8 kb, 6.3 kb, and 9.3 kb fragments from the tissue plasminogen activator (tPA) gene. For each size product, the gel shows:

Lane 1: DNA before purification Lane 2: DNA after purification by phenol/chloroform extraction Lane 3: DNA after purification with the High Pure PCR Product Purification Kit

**Result:** The High Pure Kit removed primers and primer dimers, while recovering 90% of the long PCR products (as calculated by photospectroscopy).

### **Experiment 2**

# **Preparation of hybridization probes for the analysis of number and expression patterns from sulfurtransferases from Arabidopsis**

(kindly provided by J. Papenbrock, P. von Trzebiatowski and A. Schmidt, University of Hannover, Germany)

**Background:** Sulfurtransferases are a group of enzymes widely distributed in plants, animals, and bacteria. They catalyze the transfer of a sulfane atom from a donor molecule to a thiophilic acceptor substrate.

In a database search several *Arabidopsis* EST clones with homologies to bacterial and mammalian sulfurtransferases could be identified. One of them was used to isolate a full length clone by screening an *Arabidopsis* cDNA library. The protein was expressed in *E. coli* and was determined to be a sulfurtransferase with high specificity for 3-mercapto-pyruvate. However, RNA expression data, protein contents obtained by Western blot analysis and sulfurtransferase enzyme activity measurements using 3-mercaptopyruvate or thiosulfate as substrates did not correlate with each other in *Arabidopsis* plants in different developmental states. Therefore, we hypothesize the existence of further sulfurtransferases in *Arabidopsis*. Different EST clones that show homology to sulfurtransferases and also to senescence-associated proteins are initially characterized.

**Methods:** Genomic Southern blot was performed as previously described. For preparing the hybridization probes cDNAs of interest were separated from the vectors with the respective restriction enzymes (MST1 coding for the mature 3-mercaptopyruvate sulfur-transferase, 950 bp, *Kpn* I/Pst I; EST 46D8T7, 600 bp, *Sal* I/ *Not* I). DNA fragments were separated by agarose gel electrophoresis. The gel pieces containing the MST or the 46D8T7 fragment were eluted following exactly the inserted accide fortis damg DNA from agarose gels using the High Pure PCR Product Parification for the Luted DNA was used for random prime labeling.

Northern blot was performed a previously described. The PCA probe used in Northern blotting was purified in the onlowing way. The generic coordinates was excised and the labeled frequency was purified with the high P  $\sim$  PCR Product Purification Kit according to the inserted recipe for is puting. DNA from agarose gels with an elution volume of 200  $\mu$ l.

**Results:** Hybridization of restricted genomic Arabidopsis DNA with a digoxigenin-labeled MST1 probe by random priming reveals the existence of at least two, and probably three different sulfurtransferase genes in the Arabidopsis genome (Figure 9, right). The pattern is similar to the hybridization results using the 46D8T7 cDNA but not identical (Figure 9, left).

RNA homologous to the sequence of EST clone 46D8T7 was expressed in very young plants. The expression level decreases drastically and increases continuously during aging (Figure 10). In the plants of a later developmental stage, expression level parallels the expression of SAG 13 clone, the best marker of senescence known so far.

Typical results with the kit



Figure 9: Southern blot analysis. Genomic DNA was restricted with BamH I, EcoR I, Hind III, and Xba I overnight, transferred to nylon membrane and hybridized with a digoxigenin probe labeled by random prime labeling.

Left: MST1 (3-mercapto-pyruvate sulfurtransferase) cDNA was used as a probe.

Right: cDNA from the EST clone 46D8T7 was labeled.



**Discussion:** The use of digoxigenin-labeled probes leads to very sharp bands in genomic Southern blotting. Purification of digoxigenin-labeled DNA via PCR using the High Pure PCR Product Purification Kit seems to increase the sensitivity of probes and reduce unspecific binding.

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# **High Pure RNA Isolation Kit**

for isolation of total RNA from up to 50 samples

Cat. No. 11 828 665 001

Principle	A single reagent lyses the sample and inactivates RNase. In the presence of a chaotropic salt (guanidine HCl), the released total RNA binds selectively to glass fiber fleece in a special centrifuge tube. The RNA remains bound while a DNase treatment and a series of rapid "wash-and-spin" steps remove DNA and contaminating small molecules. Finally, low salt elution removes the RNA from the glass fiber fleece. The process does	
	not require precipitation, organic solvent extractions, or extensive handling of the RNA.	
Starting material	• 10 <sup>6</sup> cultured mammalian cells (research samples)	
	• 200 – 500 μl human whole blood (research samples), containing anticoagulant	
	• $10^8$ yeast (mid- to late-log phase, $A_{600} < 2.0$ )	
	• 10 <sup>9</sup> gram positive or gram negative bacteria	
Application	• Preparation of intact total RNA, which may be used directly in RT-PCR, Differential Display RT-PCR (DDRT-PCR), Northern blotting, primer extension assays, RACE (rapid amplification of cDNA ends), cDNA library construction, <i>in vitro</i> translation, or nuclease protection assays.	
	RNA prepared from human whole blood is suitable for RT-PCR cut	
Time required	• Total time: approx. 25 min (+ sample pretreatment, if medeer)	
	• Hands-on time: <10 min (+ sample i reaction of , if needed)	
Results	<ul> <li>Yield: Variable, depending of sumple type (See the call case Part IV of "How to use the kit" in this art (1).</li> <li>Furi Printed RNA is free of Driv, protein, salts, and other cellular components.</li> </ul>	
Benefits	Saves time, 120 uS hour to an produce multiple RNA samples in minutes.	
•	<ul> <li>Ideal for a wide variety of samples, because one kit can purify RNA from many sources.</li> </ul>	
	• <b>Minimizes RNA loss</b> , because the kit removes contaminants without time-consum- ing precipitation or solvent extraction.	
	• Increases lab safety, because the kit does not use hazardous organic solvents.	

How to use the kit

2

Follow protocol Va starting with the DNase incubation (Step 5).

Vc. Isolation of total RNA from 10<sup>8</sup> yeast (S. cerevisiae)

Use fresh yeast that were harvested during mid- or late-log phase ( $A_{600} < 2.0$ ).

- 1 For each sample (10<sup>8</sup> yeast):
  - Collect the yeast by centrifugation at 2000 x g for 5 min in a standard tabletop centrifuge.
  - Resuspend the pellet in 200 µl of PBS and transfer the suspension to a sterile 1.5 ml microcentrifuge tube.

- 2 Add 10 μl lyticase solution (0.5 mg/ml) to each microcentrifuge tube. Incubate the tube for 15 min at +30°C.
- 6 Follow Protocol Va above, starting with the Lysis/Binding Buffer step (Step 2).

		Vd. Isolation of total RNA from 10 <sup>9</sup> bacteria (gram positive or gram negative)
	1	For each sample (10 <sup>9</sup> bacteria):
		Collect the bacteria by centrifugation at approximately 2000 x g for 5 min in a standard tabletop centrifuge.
		Resuspend the pellet in 200 µl of 10 mM Tris-HCl, pH 8 (, and tousier the suspension to a sterile 1.5 ml microcentrift proble.
	2	Add 4 $\mu$ I lysozyme solution (50 mc $\mu$ I $\sim$ T $<$ Hcl, pH 8.0) to each microcentrifuge tube. Incubate the tube by (0 h to $\times$ +37°C.
	3	Add 400 HL 1/S 5/ Binding Buffer to the lane mount well.
Previe	4	To transfer the sample to Highlight Tube:
		Insert on Aigh Pre-Fiter Tube into one Collection Tube.
		Protochtire sample into upper buffer reservoir of the Filter Tube.
•	5	Centrifuge the High Pure Tube assembly as in Step 4, Protocol Va.
	6	Prepare DNase solution as in Step 5, Protocol Va. After centrifugation:
		Add 100 µl DNase solution to the upper reservoir of the Filter Tube.
		Incubate the Filter Tube for 60 min at +15 to +25°C.
	7	Follow Protocol Va above, starting with the first wash (Step 6).

#### VI. Troubleshooting the High Pure protocols

The same troubleshooting procedure can be applied to all High Pure kits. For details on how to troubleshoot the above protocols, see the General Troubleshooting Procedure for all High Pure kits on page 113 of this manual. For factors that may affect the High Pure RNA Isolation Kit, see page 117.

#### Reference

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Preview from Notesale.co.uk page 61 of 260 13 The microcentrifuge tube now contains the eluted RNA. Either use 0.5 – 5 μl of the eluted RNA directly in RT-PCR or store the eluted RNA at –80°C for later analysis.

Before photometric determination of the RNA concentration, centrifuge the eluate for 2 min at maximum speed and transfer supernatant to a fresh 1.5 ml reaction tube without disturbing glass fibers at the bottom of the original tube.

### V. Troubleshooting the High Pure protocols

The same troubleshooting procedure can be applied to all High Pure kits. For details on how to troubleshoot the above protocols, see the General Troubleshooting Procedure for all High Pure kits on page 113 of this manual. For factors that may affect the High Pure FFPE RNA Micro Kit, see page 119.



### Typical results with the kit



**Figure 20: Performance of isolated RNA in RT-PCR.** RNA was isolated from a 5  $\mu$ m section of an FFPE breast tumor research sample, using either the High Pure FFPE RNA Micro Kit or a kit from another manufacturer (Supplier X). The isolated RNA samples were serially diluted and used as templates in separate RT-PCRs. A  $\beta$ -2microtubulin-specific amplification was performed, using the LightCycler<sup>®</sup> 1.5 Instrument and LightCycler<sup>®</sup> RNA Amplification Kit SYBR Green I.

**Result:** Template RNA isolated with the High Pure FFPE RNA Micro Kit performs well in the RT-PCR, giving linear results (based on the consistent slope obtained with serial dilutions) and high sensitivity (based on the early crossing points observed).

### How to use the kit

### I. Flow diagram



.

How to use the kit



### VI. Troubleshooting the High Pure Protocols

The same troubleshooting procedure can be applied to all High Pure kits. For details on how to troubleshoot the above protocol, see the General Troubleshooting Procedure for all High Pure kits on page 113 of this manual.

Typical results with the kit

### Typical results with the kit

### **Experiment 1**



**Figure 23:** The integrity and size distribution of total RNA and small RNA (2-column protocol) purified with the High Pure miRNA Isolation Kit have been checked electrophoretically on a denaturing gel (15% acrylamide/TBE/urea). A 1 μg sample of miRNA 145 was spiked into a liver tissue was before purification. Nucleic acids were visualized by ethidium bromide staining. Alternatively, electropherograms were recorded on an Agilent Bioanalyzer (1g 2 1).

Eluate volume: 100 µl; 10 µl sample per lane Eluate 1: 1-Column protocol; total RNA

**Eluate 2:** 2-Column protocol; high more la weight RNA eluted from the first filter tube (control of purification) **Eluate 3:** 2-Column protocol; burified small RNA (miRN on fine 1)

**Result** Purlieusman RNA (2-column potod I) the or DNA, nucleases and all cellular and sample containants that interfere with **P** Full The absence of contaminating DNA is examined by PCR without a preceding RT-reaction of Complific and product is obtained (data not shown).

Typical results with the kit



Figure 24: Electropherograms recorded on an Agilent Bioanalyzer Eluate 1: 1-Column protocol; total RNA

**Eluate 2:** 2-Column protocol; high molecular weight RNA eluted from the first filter tube (control of purification) **Eluate 3:** 2-Column protocol; purified small RNA (miRNA enriched) Typical result with the kit

0 The microcentrifuge tube now contains the eluted viral RNA. You may:

- EITHER use an aliquot of the eluted RNA directly in RT-PCR
  - Use 3.5 6 µl of the eluate for the reverse transcriptase reaction.
- OR store the eluted RNA for later analysis at -80°C.

### V. Troubleshooting the High Pure protocols

The same troubleshooting procedure can be applied to all High Pure kits. For details on how to troubleshoot the above protocol, see the General Troubleshooting Procedure for all High Pure kits on page 113 of this manual. For factors that may affect the High Pure Viral RNA Kit, see page 120.

### Typical result with the kit

The kit was used to prepare genomic RNA from viruses [for example, hepatitis C virus (HCV), hepatitis G virus (HGV), and human immunodeficiency virus (HIV)] for research applications. Each preparation was used as a template in RT-PCR. All these templates produced highly specific PCR products in good yield.



Figure 10. RT-FCB analysis of MS2 RI 4 iso on ed with the High Pure Urar RNA Kit. Serial dilutions of purified MS2 RNA were applied to the Filter Tubes, washed and eluted following the kit protocol. 3.5 µl of the 50 µl eluate were analyzed by two step RT-PCR using primers that resulted in a fragment of 961 bp. The indicated numbers of molecules per PCR correspond to assumed quantitative recovery.

Lane 1: 3.5 x 10<sup>7</sup> molecules/PCR; Lane 2: 3.5 x 10<sup>5</sup> molecules/PCR; Lane 3: 3.5 x 10<sup>9</sup> molecules/PCR Lane 4: 3.5 x 10<sup>2</sup> molecules/PCR Lane 5: 35 molecules/PCR Lane M: DNA Molecular Weight Marker III

**Result:** Even with theoretically 35 molecules recovered, a detectable signal in agarose gel electrophoresis is obtained.



### How to use the High Pure Extender Assembly

The High Pure Extender Assembly is delivered in single zip-bags. Five High Pure Extender Assemblies are additionally packed in labeled zip-bags. Each High Pure Extender is assembled in a 50 ml polypropylene Tube. The High Pure Extender Assembly is designed for use with table-top centrifuges and swing-bucket rotors with 4,000 x g force applicable.



#### Figure 29: High Pure Extender Assembly

Remove the High Pure Extender Assembly from the zip-bags prior to use. In the second s sample onto the g, close the High Pure Extender

Assembly unscrew the cap of the 50 ml polypropylene tube. After s m le la Assembly with the 50 ml polypropylene tube cap. After the first centrifugation step, the cample collected at the bottom of the 50 level y rot Introdyn the High Pure Extender Assembly and is ample in Stresser Forough the High Pure Extender Assembly and a v rop like rabe. Nucleic acid sare bound to the silica fleece at the urther processing carbo or the High Pure Extender Assembly from the tube containing the flow-brough. Remove the High Pure Filter Tube from After the first continue and the 50 mary for the date. Nucleic bottom of the High Pure Stranube. For unther processing end iscard the tube containing the f 50 ml polv





Figure 30: Unscrew the 50 ml polypropylene tube and remove the High Pure Extender Assembly from the 50 ml polypropylene tube.



Figure 31: Place the High Pure Extender in a new collection tube, which is placed securely in a tube rack on the bench.



.



Figure 32: Secure the High Pure Extender Assembly with one hand while grasping the locking clip (side a) of the High Pure Filter Tube cap on the opposite side with the other hand. Remove the first part of the locking clip (side a) by screwing the clip in either direction.



Figure 33: Rotate the High Pure Extender Assembly. Remove the second part of the locking clip (side b) by pulling the locking clip away from the Extender Assembly.





Figure 34: Remove the High Pure Extender from the High Pure Filter Tube by tilting the High Pure Extender away from the High Pure Filter tube toward the without the cap.



do not recomment

Pure Extended as the by. However if you decide to use a centrifuge with fixed-angle rotors, do not centrifuge High Pure Extender Assembly above  $3000 \times g$ . The use of fixed-angle rotors results in incomplete flow of the liquid through the Assembly. The remaining sample solution therefore stays within the High Pure Spin Column Filter Tube. Remove this remaining liquid by an additional spin of the High Pure Filter Tube in a bench-top centrifuge before the first washing step. For this additional spin remove the High Pure Spin Column from the High Pure Extender Assembly according to Figures 30 to 35.

Both kits are supplied with folded filters to eliminate the time-consuming centrifugation step after the alkaline lysis. In approximately 2 min (midi) or 10 min (maxi), respectively, of unattended running cellular debris and potassium dodecyl sulfate precipitates are held back by the filter thereby avoiding shearing of large DNA constructs. Besides the significant reduction of preparation time another advantage of filtration is that even small SDS precipitates which cannot be separated by conventional centrifugation are completely removed.



#### 11. **Kit contents**

- Suspension Buffer (150 ml) for suspension of bacterial cell pellets
- RNase A (15 mg) for dissolution in Suspension Buffer
- Lysis Buffer (150 ml) for bacterial cell lysis
- Neutralization Buffer (150 ml) to form a stable cellular debris precipitate
- Equilibration Buffer (70 ml) for equilibrating the columns prior to use
- Wash Buffer (370 ml) for removal of residual impurities
- Elution buffer (200 ml) for plasmid elution
- NucleoBond AX 500 Columns (10 columns) for the isolation step
- Folded filters (10 filters) to eliminate a centrifugation step and to remove cellular debris
- Sealing rings (5 rings) to station the columns in test tubes

#### **Additional materials needed** III.

- Centrifuge and tubes for harvesting bacterial cultures, capable of  $\geq$ 15,000 x g sale.co.uk
- Isopropanol
- 70% ethanol
- TE buffer or other low salt buffer
- Tube for collecting and precipitating li te
- Funnel for clearing of typa es b) folded filters PIEVIEW 143 PIEVIEW 143

Step	Ac	tion	Time / x <i>g /</i> Temperature
9		Re-insert the column into a new Collection Tube capable of withstanding high speed centrifugation ( $\geq$ 15,000 x <i>g</i> ).	
		Elute the plasmid with 15 ml Elution Buffer.	
		Allow the column to empty by gravity flow.	
•		The collected flowthrough contains the plasmid.	
U		+15 to +25°C.	
		Centrifuge immediately at high speed.	$30 \min / \ge 15,000 \ge g /$
		Carefully discard the supernatant.	+2 to +8 C
0		Wash the plasmid DNA with 4 ml chilled 70% ethanol.	+2 to +8°C
		Centrifuge at high speed.	10 min / >15,000 x g /
		Carefully remove ethanol from the tube with pipette tip.	+2 to +8°C
		Air-dry the plasmid DNA pellet.	
			10 min
12		Carefully redissolve the plasmid DNA pellet in 100 – 500 $\mu l$ TE-buffer or PCR grade water.	

### IV. Protocol for high copy number plasmid DNA, continued

	V. Troubleshooting the Genopure Plasma Vaxi protocol				
lf you get	Then, the cause may be	And you should.			
Low nucleic acid yield or purity	Kit stored to be port-optical conditions	Store kitzt + 1 to 25 Cat all times upon arrival.			
•	Buffers or other reagents were exposed to conditions that reduced their effectiveness	<ul> <li>Store all buffers at +15 to +25°C.</li> <li>After reconstitution of RNase with Suspension Buffer store aliquots at +2 to +8°C.</li> <li>Close all reagent bottles tightly after each use to preserve pH, stability and freedom from contamination.</li> </ul>			
	Reagents and samples not completely mixed	<ul> <li>Always mix the sample tube well after addition of each reagent</li> <li>Ensure Lysis Buffer and Neutralization Buffer are free of precipitates.</li> </ul>			
Low recovery of nucleic acids after elution	Non-optimal reagent has been used for elution. Salt is required for optimal elution	Use the Elution Buffer of the kit.			

3

### **DNA Isolation Kit for Cells and Tissues**

The kit offers a convenient, rapid method for the large scale isolation of intact, high molecular weight, genomic DNA from tissues, cultured cells, *E. coli*, yeast, and mouse tails.

The amount of genomic DNA recovered with the DNA Isolation Kit for Cells and Tissues is significantly higher than that obtained using alternative column-based methods in significantly less time. Starting material quantities ranging from 100 mg to 1 gram of tissue, or from  $1 \ge 10^7$  to  $5 \ge 10^7$  cultured cells, can be analyzed with the kit. Within 2 h plus resuspension time (for tissue samples), the kit yields pure genomic DNA suitable for amplification of long fragments by standard PCR, restriction enzyme digestion, and Southern blotting.

### **TriPure Isolation Reagent**

Analysis of gene expression requires clean, intact RNA templates. Isolation of intact RNA is complicated primarily by a host of stable cellular ribonucleases. To minimize RNase activity, RNA isolation procedures typically begin with cell or tissue lysis in a strongly denaturing environment. The RNA is then separated from other cellular components via multiple phenol/chloroform extractions or time-consuming CsCl step gradients.

A simpler alternative, the TriPure Isolation Reagent, offers a rapid RNA isolation procedure that can easily process multiple samples and produces 30 – 150% more RNA than other purification methods. Briefly, the reagent causes RNA, DNA, and proteil separate into different organic phases, from which each can be purifier by a series of alcohol precipitations (Chomczynski, 1993).

Thus, the TriPure Isolation Reagent allows the Sin area cous isolation of RNA, DNA, and protein from the same sample. The isolation of a could RNA is suit one to Northern blots, RT-PCR,  $poly(A^+)$  fractionation is a translation, or a Nase or the cron assays. The isolated RNA fire DNL may be used for PCR, refrection digest, or Southern blots. The denatur P on term may be analyzed on a Vector blot.



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## **DNA Isolation Kit for Mammalian Blood**

for isolation of genomic DNA from 25 samples (10 ml each)

Cat. No. 11 667 327 001

Principle	The kit procedure starts with preferential lysis of erythrocytes. The remaining leukocytes are lysed with a strong anionic detergent, and proteins are removed by dehydration and precipitation. The purified genomic DNA is then recovered via ethanol precipitation.
Starting material	• 1 – 10 ml whole blood (research samples) that has been treated with an anticoagulant (sodium heparin, sodium citrate, or EDTA)
	<i>The kit works best with fresh blood or blood stored</i> ( <i>at</i> +2 to +8°C or −15 to −25°C) for $\leq$ 3 days. It should not be used with blood that is more than 1 month old or blood that has been frozen and thawed more than 3 times.
	• Lymphocyte preparations from 10 ml mammalian blood (research samples)
	• Buffy coat preparations from 10 – 20 ml mammalian blood (research samples)
Application	• For preparation of high molecular weight, purified genomic DNA, which may be used in standard and long-template PCR, Southern blots, etc.
	The kit has been used to prepare DNA from human, moves rat, Uz, porcine, guinea pig, monkey, rabbit, and bovine blool.
Time required	• Total time: approx. 90 min (plus 10 = 0 min co resuspend DNA)
	• Hands-on time: <= mi
Results	Yield: Varia ile depending on species (or whe table under Part IV of "How to use the kit" in this article).
prev.	Purity: Avera e $\Lambda_{280}$ of isolated DNA = 1.7 – 1.9.
Benefits	• Sives time, because the kit can prepare DNA directly from whole blood without prior isolation of leukocytes or buffy coat.
	• <b>Increases lab efficiency</b> , because the kit can prepare multiple DNA samples, free of RNA and protein, in approx. 90 min.
	• <b>Increases lab safety</b> , because the kit does not require extensive handling of potentially hazardous samples, nor use hazardous organic reagents.
	• Accommodates "real world" samples with varying amounts of leukocytes, because the sample volume can be adjusted from 1 – 10 ml.

ß

How to use the kit



- Wind the DNA strands around a sterile, blunt-ended glass rod.
- Dip the glass rod containing the DNA into 3 ml cold 70% ethanol in a new sterile 50 ml centrifuge tube.
- Swirl the glass rod until the DNA strands are released into the 70% ethanol.
- Go to Step 14.
- Centrifuge the tube containing 70% ethanol and DNA for 5 min at 875 x g. Discard the 14 supernatant.
  - Dry the DNA pellet by:
    - EITHER placing the tube under vacuum without heat until the ethanol is no longer visible (usually <5 min).
    - OR inverting the tube and allowing the sample to air dry.
- To rehydrate and dissolve the DNA pellet: 16
  - Add 1 ml of Tris-EDTA (pH 8.0) (other buffer) to the tube.
  - Vortex thoroughly.
    - Incubate at +65°C for 30 60 min.
      - For DNA from human blood, use a 30 min incubation. For DNA from other mammalian species, use a 60 min incubation.
  - During the incubation, periodically vortex the tubes to facilitate solubilization of the DNA.
    - Do not incubate the tube for >60 min at  $+65^{\circ}C$ . Store DNA at +2 to  $+8^{\circ}C$ .
  - Store DNA at +2 to +8°C

### Isol otocol the DNA

you should... If you get cal cloudy upper Start with fresh samples and repeat Steps 1 – 4 of the protocol laver and a red red cell lysis with one or more changes: lower layer at Be sure blood is at +15 to +25°C before starting protocol. Step 4 Use a 15 min incubation in Step 3. ▶ If mixing by hand in Step 3, invert the tube more frequently. Particles present Incomplete Using fresh samples, repeat Protocol Va, with one or more changes: in sample after white cell lysis In Step 6, be sure the white cell pellet is fully resuspended in vortexing the residual supernatant after vortexing. (Step 7) After adding buffer in Step 7, vortex the sample until the solution is dark brown/red and contains no particles. In Step 7, add enough White Blood Cell Lysis Buffer to ensure the solution is not viscous and does not contain clumps of cells. In Step 7, incubate the tube at  $+37^{\circ}$ C for 15 - 30 min to facilitate lysis.

Typical results with the kit

### Typical results with the kit

### **Experiment 1**



Figure 46: Use of DNA from human and mouse blood (prepared with the DNA Isolation Kit for Mammalian Blood) as templates for the amplification of large DNA fragments. Human and mouse blood research samples were collected as previously described (Noeth and Dasovich-Moody, 1997). DNA was prepared from 10 ml of each sample according to the protocol. Aliquots of each DNA preparation were used as templates for the long-template PCR amplification of several gene fragments. Each amplification procedure was performed with the Expand Long Template PCR System, using instructions given in the package insert.

Left panel shows gene fragments amplified from human DNA:

Lanes 1, 10: DNA Molecular Weight Marker III Lanes 2, 3: tPA fragment (9.3 kb) amplified from 25 ng DNA

Lanes 4, 5: tPA fragment (15 kb) amplified from 50 ng DNA

**Lanes 6, 7:**  $\beta$ -globin fragment (23 kb) amplified (20 kb) amplified (20 kb) amplified (20 kb) amplified from **Lanes 8, 9:**  $\beta$ -globin fragment (28 kb) amplified from

Right panel shows the following fragments amplified from mouse DNA:

**Lanes 1, 10:** DNA Molecular Weitht Marker III **Lanes 2, 3:** IL-2 gene (4.2 kH and n d from 330 ng DNA **Lane:** 4  $\odot \alpha$  -2 rollagen fragment (5.6 kb) amplified

f.<sup>pp</sup>, h0 sc DNA Lones **6, 7:** α-2 collagen fragment (10.4 kb) amplified from 50 ng DNA Lone 5.2, to 2 collagen fragment (15.4 kb) a...plk = from 100 ng DNA

Experiments were clarity is in gels.

2 3 4

1

200 ng DN

Figure 47: Use of DNA research samples (pre Isolation Kit for Mamm of the n-ras gene by Stabove protocol was used several human blood sar been prepared with a dift was also prepared from a and a buffy coat prepara the protocol outlined in t of each preparation was electrophoretically separ membrane. DNA on the a DIG-labeled n-ras protochemiluminescently. The Lanes 1, 12: DNA Mole Lanes 2, 3: whole blood lant Lanes 4, 5: whole blood lant

Figure 47: Use of DNA from various human blood research samples (prepared with the DNA Isolation Kit for Mammalian Blood) for detection of the n-ras gene by Southern hybridization. The above protocol was used to prepare DNA from several human blood samples, each of which had been prepared with a different anticoagulant. DNA was also prepared from a lymphocyte preparation and a buffy coat preparation, using a modification of the protocol outlined in the kit package insert. Ten µg of each preparation was digested with Eco RI, electrophoretically separated, and blotted to a nylon membrane. DNA on the membrane was hybridized to a DIG-labeled n-ras probe, and the results visualized chemiluminescently. The samples used were: Lanes 1, 12: DNA Molecular Weight Marker VII Lanes 2, 3: whole blood, sodium citrate anticoagu-

Lanes 4, 5: whole blood, heparin anticoagulant Lanes 6, 7: whole blood, sodium EDTA anticoagulant Lanes 8, 9: buffy coat preparation Lanes 10, 11: lymphocyte preparation

**Result:** Each lane contained only a single hybridization band of the expected (7.2 kb) size (Taparowsky et al., 1983).





How to use the reagent



# **mRNA** Isolation Kit

for the isolation of poly(A)+ RNA (at least 70  $\mu g)$ 

Cat. No. 11 741 985 001

Principle	The poly(A) <sup>+</sup> tail of mRNA hybridizes to a biotin-labeled oligo(dT) probe. Streptavidin- coated magnetic particles capture the biotinylated dT-A hybrids. A magnetic particle separator collects the magnetic particles, and washes remove contaminants. Then, water elutes the mRNA from the particles.				
Starting material	Research samples may contain up to:				
	• 2.5 mg total RNA				
	• 10 <sup>8</sup> cultured cells				
	• 1 g tissue				
Application	Preparation of highly purified poly(A) <sup>+</sup> RNA, which may be used for RT-PCR, cDNA synthesis, Northern blotting, Northern ELISA, RNase protection assays, <i>in vitro</i> translation, etc.				
Time required	• Total time: approx. 30 min (starting from total RNA)				
	• Hands-on time: approx. 15 min (starting from total RNA)				
Results	• Yield: Variable, depending upon starting material (See Part IV of "How to use the kit in this article).				
	• Purity: Isolated mRNA, free of DNA and other a las				
Benefits	• Saves time, because the kit can iso an n.ENA directly from a llysates and tissue homogenates; no isol and of total RNA required.				
	• Account divites a wide range of samples, and allows both small and large scale				
Pr	<b>Increases D</b> safe was cause the kit does not use hazardous organic solvents.				

5

### How to use the kit

### I. Flow diagram



	mkina isolation Kit		
	How to use the kit		
			Cultured cells: Wash cells (2 x 10 <sup>7</sup> ) twice with ice-cold phosphate-buffered saline (PBS). Add 3 ml Lysis Buffer to the cell pellet. Shear DNA mechanically by passing the sample 6 x through a 21-gauge needle.
			Tissue: Snap freeze 200 mg of tissue. Grind frozen tissue to a homogeneous powder in a pre-cooled mortar. Chill 3 ml Lysis Buffer to $\leq 0^{\circ}$ C in a sodium chloride-ice water bath, then add the frozen powder to the chilled Lysis Buffer. Homogenize the powder suspension by passing it 4 x through a 21-gauge needle. Centrifuge the suspension at 11,000 x g for 30 s. Use only the supernatant for the capture procedure.
			0°C and -4°C.
		2	Prepare the Streptavidin-coated Magnetic Particles:
			Resuspend Streptavidin-coated Magnetic Particles thoroughly and pipette 300 µl Streptavidin-coated Magnetic Particles into a fresh cup or tube.
			Immobilize the Streptavidin-coated Magnetic Particles on the side of the container with a Magnetic Particle Separator.
			Remove the storage buffer from the Streptavidin-coated Magnetic Particles.
			Resuspend the Streptavidin-coated Magnetic Particles in 500 µl Lysis Buffer.
			Again immobilize the Streptavidin-coated Magnetic Particles with the magnetic separator and remove all the Lysis Buffer.
			Never let the Streptavidin-coated Magnetic Particles dry out.
		3	Hybridize the mRNA to the Biotin-labeled $Oligo(dT)_{20}$ :
5			<ul> <li>Add 3 µl Biotin-labeled Oligo(dT)<sub>20</sub> to the sample (from Step 1)</li> <li>Mix sample and Oligo(dT)<sub>20</sub> well to form the reprintization mix.</li> </ul>
	, ie	4	<ul> <li>Immobilize the biotinylated JCA is price with Streptavidin-coated Magnetic Particles:</li> <li>Add hybridization must one tube containing the prepared Streptavidin-coated</li> <li>Magnetic Particles (from Step 2):</li> <li>Resuspend Streptavion-poated Magnetic Particles in the hybridization mix.</li> </ul>
	Previ		<ul> <li>Incubation:</li> <li>Incubati</li></ul>
		-	Separate the Streptovidin, costed Magnetic Particles from the fluid with a Magnetic
		0	Particle Separator.
			This step takes about 3 min.
		6	Wash the Streptavidin-coated Magnetic Particles 3 times. For each wash:
			Add 500 μl Wash Buffer to the tube.
			Resuspend the Streptavidin-coated Magnetic Particles in the Wash Buffer.
			<ul> <li>Separate the Streptavidin-coated Magnetic Particles from the buffer with a Magnetic Particle Separator.</li> </ul>
			Remove all Wash Buffer and discard.
		7	Elute the mRNA:
			Add 50 μl redistilled water to the Streptavidin-coated Magnetic Particles.
			Resuspend the Streptavidin-coated Magnetic Particles in the redistilled water.
			Incubate for 2 min at +65°C.
			Separate Streptavidin-coated Magnetic Particles from the eluate with a Magnetic Particle Separator.
		8	Transfer the supernatant (containing the mRNA) to a fresh RNase-free tube.
			<i>I</i> To quantitate the mRNA spectrophotometrically, assume that $1 A_{260}$ unit corresponds to a concentration of 40 $\mu$ g/ml.

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		Protocol for purifying radiolabeled nucleic acids	
		Remove the column from the storage bag, and gently invert it several times to resuspend the matrix.	•
	2	Remove most of the buffer from the column as follows:	
		Remove the top cap from the column, then remove the bottom tip.	
		Drain the column by gravity, then discard the eluate.	
		Remove residual buffer from the column as follows:	
		Place the column in a collection tube.	
		Centrifuge at 1100 x g for 2 min in a swinging-bucket rotor.	
		Discard the collection tube with the eluted buffer.	
	4	While keeping the column upright, very carefully apply the sample to the center of the column bed.	
		Apply 20 – 50 μl sample to a G-25 column; 20 – 100 μl to a G-50 column.	
	5	While keeping the column upright, place the column in the second collection tube.	
	6	Centrifuge the tube at 1100 x $g$ for 4 min in a swinging-bucket rotor.	
	7	Save the eluate from the second collection tube. It contains the purified sample.	
	8	Discard the column in a radioactive waste container.	
	V.	Troubleshooting the Quick Spin portscol	
If you get	V. Then caus	Troubleshooting the Quick Spin portecol	
<b>If you get</b> Poor sample recovery	V. Then caus Colur caus sam nucle recov	Troubleshooting the Quick Spin portecol , the angree ng of the nn, which is such frequence te and reduced te acids very	
If you get Poor sample recovery	V. Then caus Colur caus sam nucle recov Wror	Ind poissould       Constrained         Ind poissould       100         Ing of the nn, which is pace flawood le and reduced the and	-
If you get Poor sample recovery	V. Then caus Colur caus sam nucle recov Wror durir tion	Troubleshooting the Quick Spin parts col         , the enapped         ing of the nn, which especially after centrifugation.         Space fawori le and reduced ic acids very         ng g-force used ng centrifuga-         • Use 1100 x g for centrifugation spins.         • Be sure the centrifuge is correctly calibrated.	· ·
If you get Poor sample recovery	V. Then caus Colur caus sam nucle recov Wror durin tion Wror	Troubleshooting the Quick Spin portecol         , the enaybe:         enaybe:         ng of the nn, which spin the column upright during and after application of cample, especially after centrifugation.         Space faworie         le and reduced tic acids very         ng g-force used ag centrifuga-         le sure the centrifuga is correctly calibrated.         le sure the centrifuge is correctly calibrated.         ng rotor	- -
If you get Poor sample recovery	V. Then caus Colur caus sam nucle recov Wror durin tion Wror	Troubleshooting the Quick Spin pactecol         , the enables       Ind pace involut         and of the number of the	· · ·
If you get Poor sample recovery	V. Then cause Colur cause sam nucle recov Wror durir tion Wror durir tion Wror	Troubleshooting the Quick Spin Pattecol         , the engyber         ng of the nn, which spice thwo le and reduced cic acids very         ng g-force used ng centrifuga-         >         Use 1100 x g for centrifugation spins.         >         >         >         ng rotor         >         Nuch or too DNA/RNA ed on column         DNA/RNA         >         Apply a sample containing 0.02 – 1.0 mg/ml nucleic acid. At >1.0 mg/ml, the sample is viscous and may not migrate through the columns easily, leading to poor recovery and/or contamination with smaller molecules. At <0.02 mg/ml, DNA/RNA with recovery may be low.	· · ·

Typical result with the columns

### Typical result with the columns



Figure 58: Removal of Unincorporated Radiolabeled Nucleotides from 5' End-labeled Oligomers using the Mini Quick Spin DNA and Oligo Columns. Various oligomers, ranging in size from 32 bp to 8 bp, were end-labeled using the 5' End-Labeling Kit and  $\gamma^{32}$ P-ATP and purified using either the mini Quick Spin DNA or Oligo Columns to remove unincorporated nucleotide, per manufacturer's instructions. One fifth of the isolated sample was electrophoresed through a 20% acrylamide gel in 1x TBE for 5 - 6 hours at 1501/ e gel was overlaid with plastic wrap and exposed to Lumi-Film for approximately 18 h.



prev

1e.co Lane 2: Oligo Marker/miniQuick Spin DNA Column Lane 3: Oligo Marker/mini Quick Spin DNA Lane 4: -Lane 5: Oligo Marker Lane 6: Oldo ni Quick Spin Oligo Columr mini Quick Spin 🕯 nigo Cr Lane um ane 9: Oligo Marke nincorporated Radio-labeled Nucleotides from 5' End-labeled Oligomers using the Mini pin DNA and Mini Quick Spin Oligo Columns Ouicl

### References

Lane 1: Oligo Marker Control

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# Automated Nucleic Acid Isolation

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### **Real-Time PCR Instruments Available from Roche Applied Science**

Real-time PCR offers an alternative method for both qualitative and quantitative analysis. This type of analysis allows the amplification and fluorescent detection steps to be performed by a single instrument in a single tube with data recorded online. A real-time PCR instrument measures the accumulation of PCR products during amplification with fluorescent dyes. Because PCR itself and the detection of PCR products occur in the same reaction (vessel), this set-up is also called "homogeneous PCR". The LightCycler® System incorporates several features that make it the ideal tool for qualitative and quantitative PCR as well as mutation analysis in general laboratory applications. It includes instrumentation, software, reagents, technical support, and application-specific kits.

Two real-time PCR systems are available from Roche Applied Science:



**The LightCycler® Carousel-Based System** (LightCycler® 2.0 Instrument, Cat. No.: 03 531 414 201).

The LightCycler<sup>®</sup> 2.0 Instrument is optimized for two fluorescence detection formats: SYBR Green I and HybProbe probes. In addition, the instrument supports a wide variety of other fluorescence detection formats, such as monocolor SimpleProbe probes, hydrolysis probes, and other formats based on FRET (fluorescence resonance energy transfer).



**The LightCycler**<sup>®</sup> **480 System** (Cat. No  $0.4642265001 \rightarrow 96$  well; Cat. No. 04 545 885 001  $\rightarrow$  384 cell is a modular online PCR device for qualitative or cualitianty to 25 ccon of nucleic acids, mutation screening and reactly ling. It meets the sector of broad range of contributions in genomic research, such as array validation, gene-knockdown scales and SUP analysis.

Offering the sending trand accuracy one has come to expect only from R and Applied Science's LightCycler® Carousel-Based System, to LightCycler® 480 Real-time PCR System goes one step further providing enhanced 96- or 384-multiwell throughput.

The LightCycler® 480 System setup enables the use of all current probe formats (*e.g.*, SYBR Green I, HybProbes probes, SimpleProbe probes, hydrolysis probes and ResoLight dye for high resolution melting).

For even higher throughputs (up to 1536 reactions in one run) and smaller reaction volumes (0,5-2 $\mu$ l), a separate plate-based instrument, the LightCycler® 1536 System, is available.

For details concerning the instruments, reagents and software please refer to http://www.lightcycler.com, http://www.lightcycler480.com and www.lightcycler1536.com



ResoLight is a trademark of Roche

# **Ordering Guide**

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### **MagNA Lyser Instrument and Accessories**

Product	Application	Cat. No.	Pack Size
MagNA Lyser	Automated homogenization of tissue samples	03 358 968 001	1 instrument
Instrument <sup>#</sup>		03 358 976 001	plus accessories
MagNA Lyser	Holds up to 16 sample tubes for tissue homogenization	03 359 093 001	1 rotor
Rotor <sup>#</sup>			
MagNA Lyser	Houses the MagNA Lyser Rotor to maintain the	03 359 085 001	1 cooling block
Rotor Cooling	temperature of the samples at +2 to +8°C		
Block <sup>#</sup>			
MagNA Lyser	Specially designed ceramic beads to achieve optimal	03 358 941 001	100 tubes
Green Beads <sup>#</sup>	homogenization of various sample materials		(prefilled with ceramic beads)

### Automated Isolation using the MagNA Pure LC Instrument

Product	Application	Cat. No.	Pack Size
MagNA Pure LC Instrument <sup>+</sup>	Robotic workstation for fully automated nucleic acid preparation and filling of LightCycler <sup>®</sup> Capillaries, 96-well PCR plates, and tubes suitable for the most commonly used PCR instruments	12 236 931 001	1 instrument plus accessories
<b>Reagent Kits for Iso</b>	lation of genomic DNA		
MagNA Pure LC	Ready-to-use reagents for the isolation of high-quality	03 003 999 001	H C
<b>DNA</b> Isolation	genomic DNA from whole blood, white blood cells,		(192 reactions)
Kit I⁺	peripheral blood lymphocytes, and cultured cells, o in a	5.0	
	the MagNA Pure LC Instrument.		
MagNA Pure LC	Ready-to-use reagents for the hid at on of high-quality.	0. 186 229 001	1 kit
DNA Isolation	intact genomic EV. A rom a wide variety of rumar a d		(192 reactions)
Kit II <sup>+</sup> (Tissue)	animal triste suppres, using the Magin A trie Lu		
MogNA Pro.C	Ready-to-use reace its for the isolation of high-quality,	03 264 785 001	1 kit
I IA solution	inta Dav. n. Unnungal DNA from the most difficult-		(192 reactions)
Kit III+ (Bacteria &	to-p ocess research sample materials, using the		
Fungi)	MagNA Pure LC Instrument.		
MagNA Pure LC	Ready-to-use reagents for the purification of genomic	03 310 515 001	1 kit
DNA Isolation Kit*	DNA from large amounts of whole blood (up to 1 ml),		(96 – 288
– Large Volume	blood cells, or culture cells (up to 5 x 10 <sup>6</sup> ), using the		reactions)
	MagNA Pure LC Instrument.		
Reagent Kits for Iso	lation of viral and total Nucleic Acids		
MagNA Pure LC	Ready-to-use reagents for the purification of viral	03 038 505 001	1 kit
Total NA Isolation	nucleic acid from serum, plasma and whole blood,		(192 reactions)
Kit <sup>*</sup>	Using the MagNA Pure LC Instrument	00.00/ 700.001	4.1.1
MagNA Pure LC	Ready-to-use reagents for the purification of viral	03 264 793 001	I KIL
	and pleased using the MegNA Pure I C Instrument		(192 reactions)
Volume	and plasma, using the MagNA Pure LC instrument		
<b>Reagent Kits for Iso</b>	lation of total RNA		
MagNA Pure LC	Ready-to-use reagents, developed to maximize yield of	03 542 394 001	1 kit
<b>RNA Isolation Kit+-</b>	purified total RNA with superior quality isolated from		(192 reactions)
High Performance	blood, blood cells or culture cells, using the MagNA		
	Pure LC Instrument		

\* CE marked /for USA for laboratory use

+ For general laboratory use.

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Features	Cat. No.	Pack Size
<ul> <li>Rapidly and completely denatures most proteins</li> <li>Particularly suited for gel electrophoresis</li> </ul>	11 667 289 001	1 kg
Virtually free of proteases, RNases, and DNases	10 708 976 001	1 ka
► pK 8.3 (at +20°C), which makes Tris ideal for preparing buffers in the biological	11 814 273 001	5 kg
pH range		
Used in many published purification procedures		
► Ultrapure	11 332 481 001	5 x 10 ml

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#### Disclaimer

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