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Qproteome™ Nuclear Protein Handbook

For isolation of nuclear and nucleic-acid
binding proteins from eukaryotic cell lysates



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Kit Contents	4
Storage	4
Product Use Limitations	4
Quality Control	4
Product Warranty and Satisfaction Guarantee	5
Technical Assistance	5
Safety Information	5
Introduction	6
Fractionation of Cell Lysate	6
Principle and procedure	7
Preparation of the cytosolic fraction	8
Preparation of the nucleic-acid binding protein fraction	8
Preparation of the “insoluble” nuclear protein fraction	8
Protocols	9
■ Isolation of Nuclear Proteins from Mammalian-Cell Lysates	9
■ Acetone Precipitation of Protein Fractions	11
Troubleshooting Guide	12
Ordering Information	13
QIAGEN Distributors	15

Kit Contents

Qproteome Nuclear Protein Kit	
Catalog no.	37582
Number of preps	12
Lysis Buffer NL	2 x 10 ml
Extraction Buffer NX1	1 ml
Extraction Buffer NX2	2 x 1 ml
Detergent Solution NP	2 x 0.25 ml
DTT Stock Solution	0.25 ml of a 1 M solution
Benzonase [®]	2000 Units (25 U/ μ l)
Protease Inhibitor Solution (100x)	300 μ l
Handbook	1

Storage

Buffers NL, NX1, and NX2 should be stored at room temperature (15–25°C).

Detergent Solution NP and Protease Inhibitor Solution (100x) should be stored at 4°C.

Benzonase[®] and DTT Stock Solution should be stored at –20°C.

Product Use Limitations

Qproteome Nuclear Protein Kits are developed, designed, and sold for research purposes only. They are not to be used for human diagnostic or drug purposes or to be administered to humans unless expressly cleared for that purpose by the Food and Drug Administration in the USA or the appropriate regulatory authorities in the country of use. All due care and attention should be exercised in the handling of many of the materials described in this text.

Quality Control

In accordance with QIAGEN's ISO-certified Total Quality Management System, each lot of QIAamp MinElute Media Kit is tested against predetermined specifications to ensure consistent product quality.

Product Warranty and Satisfaction Guarantee

QIAGEN guarantees the performance of all products in the manner described in our product literature. The purchaser must determine the suitability of the product for its particular use. Should any product fail to perform satisfactorily due to any reason other than misuse, QIAGEN will replace it free of charge or refund the purchase price. We reserve the right to change, alter, or modify any product to enhance its performance and design. If a QIAGEN product does not meet your expectations, simply call your local Technical Service Department or distributor. We will credit your account or exchange the product — as you wish.

A copy of QIAGEN terms and conditions can be obtained on request, and is also provided on the back of our invoices. If you have questions about product specifications or performance, please call QIAGEN Technical Services or your local distributor (see inside back cover).

Technical Assistance

At QIAGEN we pride ourselves on the quality and availability of our technical support. Our Technical Service Departments are staffed by experienced scientists with extensive practical and theoretical expertise in molecular biology and the use of QIAGEN® products. If you have any questions or experience any difficulties regarding Qproteome Nuclear Protein Kits or QIAGEN products in general, please do not hesitate to contact us.

QIAGEN customers are a major source of information regarding advanced or specialized uses of our products. This information is helpful to other scientists as well as to the researchers at QIAGEN. We therefore encourage you to contact us if you have any suggestions about product performance or new applications and techniques.

For technical assistance and more information please call one of the QIAGEN Technical Service Departments or local distributors (see inside back cover).

Safety Information

When working with chemicals, always wear a suitable lab coat, disposable gloves, and protective goggles. For more information, please consult the appropriate material safety data sheets (MSDSs). These are available online in convenient and compact PDF format at www.qiagen.com/ts/msds.asp where you can find, view, and print the MSDS for each QIAGEN kit and kit component.

24-hour emergency information

Emergency medical information in English, French, and German can be obtained 24 hours a day from:

Poison Information Center Mainz, Germany

Tel: +49-6131-19240

Introduction

The functional architecture of the eukaryotic cell nucleus is of great interest to cell biologists. The identification of nuclear proteins — especially nucleic-acid-binding proteins (e.g., transcription factors) — is important for an understanding of genome regulation and function, and provides clues about the molecular function of novel proteins.

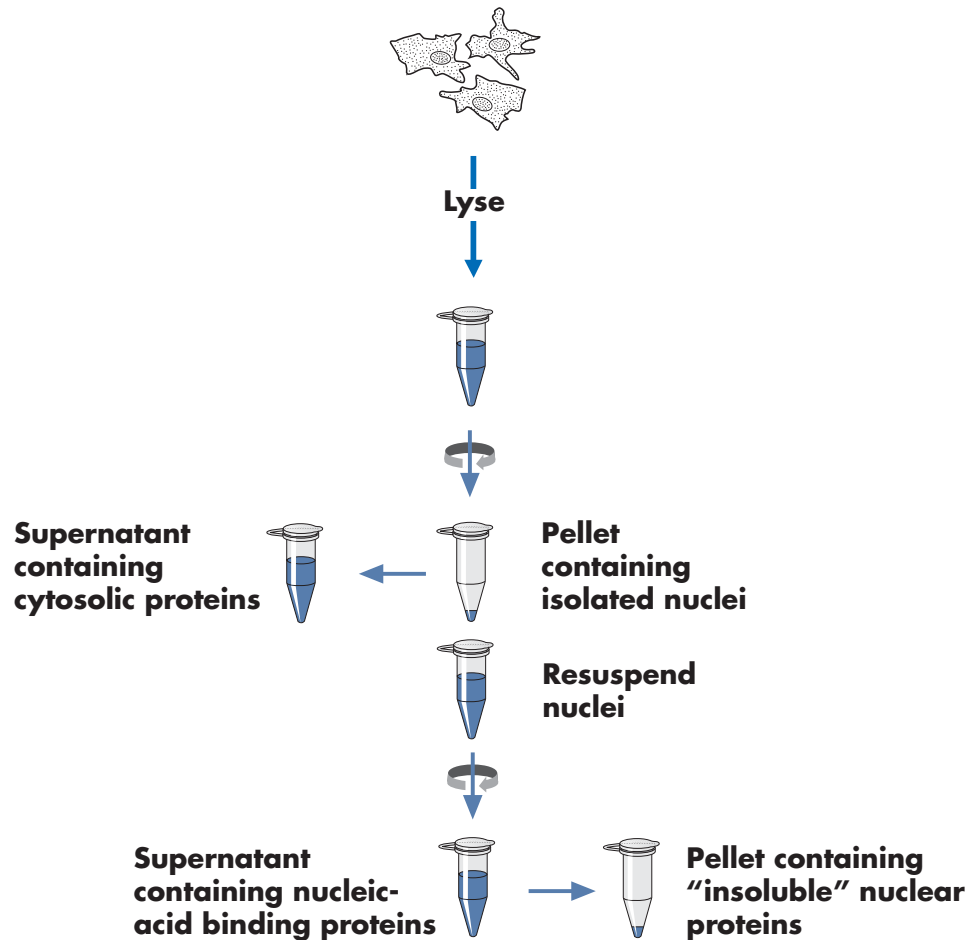
The nucleus contains a cell's genetic information and is the site of gene expression. Biological processes involving nucleic acids, such as transcription, replication, recombination, and DNA repair all involve the action of proteins that bind nucleic acids in a sequence-specific manner. These proteins interact with other proteins, which may or may not bind directly to nucleic acids. As a result of these interactions, large functional complexes are formed and anchored to specific nucleic acid sites. Many auxiliary proteins, enzymes, and complexes in the nucleus have important, general functions and are present in small to moderate amounts. However, proteins that bind at selected sites, such as transcription factors bound to specific promoters, represent only 0.01–0.001% of total cellular protein.

Fractionation of Cell Lysate



Figure 1 Fractions obtained after processing cell lysates using the Qproteome Nuclear Protein Kit. **C:** Cytosolic fraction; **D:** nucleic-acid-binding protein fraction; **I:** “Insoluble” fraction.

Nuclear Protein Fractionation Procedure



Principle and procedure

The Qproteome Nuclear Protein Kit is designed for specific enrichment of nuclear proteins. The kit provides standardized sample preparation for reliable quantitative and qualitative analysis of proteins for targeted proteomics. Nuclear proteins are separated from complex protein mixtures, allowing detection of low-abundance proteins (e.g., most transcription factors). Starting material for one fractionation procedure is 5×10^6 – 1×10^7 cells. The procedure has been used successfully with several different mammalian cell lines including HeLa, Jurkat, NIH3T3, HEK293, and Cos (Table 1, page 8). QIAGEN also offers the Qproteome Nuclear Subfractionation Kit (cat. no. 37531) which provides further subfractionation of the nucleic acid binding protein fraction.

Table 1. Typical Total Protein Yields in Cytosolic, DNA Binding Protein, and Histone Fractions

Cell type	Number of cells processed	Cytosolic fraction	Nucleic-acid-binding proteins	Histones
HeLa	5.4×10^6	1.9 mg	0.34 mg	0.40 mg
HEK293	4.0×10^7	9.4 mg	0.55 mg	0.89 mg
Jurkat	1.5×10^7	2.6 mg	0.23 mg	0.53 mg
Cos	3.7×10^6	2.9 mg	0.14 mg	0.18 mg
NIH3T3	7.9×10^6	1.9 mg	0.28 mg	0.34 mg

Preparation of the cytosolic fraction

Cells are incubated in hypotonic buffer, causing them to swell. Detergent added to the lysis buffer ruptures the plasma membrane and centrifugation is used to separate the cytosolic fraction (supernatant) from the cell nuclei (pellet).

Preparation of the nucleic-acid binding protein fraction

The nuclear pellet is washed to remove cytosolic contaminants from the isolated cell nuclei. Washed cell nuclei are incubated in a buffer containing a high concentration of salt. During this incubation step, the nuclei shrink and nucleic-acid-binding proteins (e.g., transcription factors) separate from nucleic acids and diffuse through the nuclear pore to the exterior of the nucleus. A centrifugation step separates the nucleic-acid-binding proteins (which are found in the supernatant) from the nuclear debris, which includes genomic DNA (the pellet). The expected yield from one fractionation procedure is 200–300 μg specific nucleic-acid-binding proteins, at a concentration of approximately 5 $\mu\text{g}/\mu\text{l}$. This nucleic-acid-binding protein fraction can be used for activity tests (e.g., gel-shift assays, transcription factor activity assays) and characterization of function without further processing.

Preparation of the “insoluble” nuclear protein fraction

Extraction of “insoluble” nuclear proteins is performed by incubation of the nuclear debris with a buffer containing Benzonase[®], a DNase/RNase which digests genomic DNA and releases nuclear proteins intimately associated with DNA (e.g., histones).

Protocol: Isolation of Nuclear Proteins from Mammalian-Cell Lysates

Equipment and reagents to be supplied by the user

- Ice-cold PBS
- Cell scraper
- 15 ml conical tube
- Microcentrifuge tubes
- Thermomixer or similar
- Optional: Acetone stored at -20°C

Important notes before starting

- All steps are performed at 4°C . Use pre-cooled buffers and equipment. Separated protein fractions should be stored at -80°C .
- For downstream applications, such as 2-D gel analysis, fractions may need to be concentrated. This can be achieved by acetone precipitation (see page 11).
- Starting material for one fractionation procedure using the Nuclear Protein Kit protocol is $5 \times 10^6 - 1 \times 10^7$ cells.

Things to do before starting

- Dilute $10 \mu\text{l}$ of the 1 M DTT Stock solution with $90 \mu\text{l}$ deionized, sterile water to give a concentration of 0.1 M. Store 0.1 M DTT Solution at -20°C .
- Immediately before starting the protocol, supplement the buffers used in the protocol with Protease Inhibitor Solution, DTT Stock Solution, and Benzonase[®] as shown in Table 2.

Table 2. Preparing Buffers for the Qproteome Nuclear Protein Kit Protocol

Buffer	Required volume per prep.	Protease Inhibitor Solution	0.1 M DTT	Benzonase [®]
Lysis Buffer NL (for protocol steps 6 and 11)	$1000 \mu\text{l}$	$10 \mu\text{l}$	$5 \mu\text{l}$	–
Extraction Buffer NX1 (step 14)	$50 \mu\text{l}$	$0.5 \mu\text{l}$	–	–
Extraction Buffer NX2 (step 17)	$100 \mu\text{l}$	$1 \mu\text{l}$	$1 \mu\text{l}$	$1 \mu\text{l}$

Procedure

Cell collection

1. Aspirate cell-culture medium from culture plate.
2. Wash cells twice with 5 ml ice-cold PBS.
3. Add 10 ml ice-cold PBS.
4. Remove cells from culture plate by gentle scraping with cell-scraper and transfer cells to a pre-chilled 15 ml conical tube.
5. Centrifuge cell suspension for 5 min at 450 x g in a centrifuge pre-cooled to 4°C. Discard supernatant. Keep cell pellet on ice.

Cell lysis and isolation of cell nuclei

6. Gently resuspend the cells collected in step 5 in 500 μ l Lysis Buffer NL (supplemented with Protease Inhibitor Solution and 0.1 M DTT, see Table 2) by pipetting up and down several times. Incubate for 15 min on ice.
7. Transfer the resuspended cells to a clean pre-chilled microcentrifuge tube.
8. Add 25 μ l Detergent Solution NP to the cell suspension and vortex for 10 s at maximum speed.
9. Centrifuge cell suspension for 5 min at 10,000 x g in a microcentrifuge pre-cooled to 4°C.
10. Transfer the supernatant (cytosolic fraction) into a new microcentrifuge tube and store at -80°C.
11. Resuspend the pellet (which contains cell nuclei) in 500 μ l Nuclear Protein Lysis Buffer NL (supplemented with Protease Inhibitor Solution and 0.1 M DTT, see Table 2) by vortexing for 5 s at maximum speed.
12. Centrifuge the suspension of nuclei for 5 min at 10,000 x g in a pre-cooled microcentrifuge at 4°C.
13. Discard the supernatant and save the nuclear pellet.

Extraction of nucleic-acid-binding proteins

14. Resuspend the nuclear pellet from step 13 in 50 μ l Extraction Buffer NX1 (supplemented with Protease Inhibitor Solution, see Table 2) by pipetting up and down. Incubate suspension for 30 min with gentle agitation (e.g., 750 rpm in a Thermomixer) at 4°C.
15. Centrifuge the suspension of nuclei for 10 min at 12,000 x g in a microcentrifuge pre-cooled to 4°C.

16. Transfer the supernatant (which contains nucleic-acid binding proteins) into a new microcentrifuge tube.

Store the supernatant and the pellet at -80°C until you are ready to proceed with further analysis. This fraction can be used directly for activity assays (e.g., gel-shift assays).

Continue with step 17 to use the pellet for extraction of “insoluble” nuclear proteins (e.g., histones).

Extraction of “insoluble” nuclear proteins (e.g., histones)

- 17. Resuspend the pellet from step 16 in 100 μl Extraction Buffer NX2 (supplemented with Benzonase[®], Protease Inhibitor Solution, and 0.1 M DTT Stock Solution, see Table 2) by pipetting up and down. Incubate suspension for 1 h with gentle agitation (e.g., 750 rpm in a Thermomixer) at 4°C .**
- 18. Centrifuge pellet suspension for 10 min at 12,000 x g in a pre-cooled microcentrifuge at 4°C . Transfer the supernatant to a new microcentrifuge tube and store at -80°C .**

Protocol: Acetone Precipitation of Protein Fractions

This protocol is suitable for concentrating and desalting protein samples for downstream applications such as SDS-PAGE.

- 1. Add four volumes of ice-cold acetone to the protein fraction and incubate for 15 min on ice.**
- 2. Centrifuge for 10 min at 12,000 x g in a pre-cooled microcentrifuge at 4°C . Discard the supernatant and air dry the pellet.**
Do not overdry the pellet as this may make it difficult to resuspend.
- 3. Depending on the application, resuspend the pellet in the required sample buffer.**

Troubleshooting Guide

This troubleshooting guide may be helpful in solving any problems that may arise. The scientists in QIAGEN Technical Services are always happy to answer any questions you may have about either the information and protocol in this handbook or molecular biology applications (see back page for contact information).

Comments and suggestions

Low protein concentration in protein fractions

Too few cells in starting material Use a minimum of 5×10^6 cells per preparation (see Table 1).

Poor protein compartmentalization

a) Incomplete removal of cytosolic fraction Ensure that all traces of supernatant are removed after the nuclear pellet wash step.

b) Nuclei are disrupted during nucleic-acid-binding protein extraction Gently resuspend the nuclear pellet in extraction buffer NX1 by pipetting up and down. Do not vortex.

No or low protein activity

Fractionated proteins are degraded Ensure all buffers and equipment are cooled to 4°C during the entire procedure. Ensure that Protease Inhibitor Solution has been added to buffers. Snap-freeze eluted proteins using liquid nitrogen.

Ordering Information

Product	Contents	Cat. no.
Qproteome Nuclear Protein Kit	For 12 nuclear protein preparations: Buffers, Reagents, Protease Inhibitor Solution, Benzonase [®]	37582
Related products		
Qproteome Nuclear Subfractionation Kit	For 6 nuclear protein preparations: Buffers, Reagents, Nuclear protein Fractionation Columns (6), Nuclear Protein Fractionation Resin, Protease Inhibitor Solution, Benzonase [®]	37531
Qproteome Albumin/IgG Depletion Kit	For albumin/IgG depletion of 6 serum or plasma samples: Albumin/IgG Depletion Spin Columns (6)	37521
Qproteome Murine Albumin Depletion Kit	For albumin depletion of 6 murine serum or plasma samples: Murine Albumin Depletion Spin Columns (6)	37591
Qproteome Bacterial Protein Prep Kit	For soluble protein preparations from up to 4.5 liters of bacterial cell culture: Lysis Buffer, Lysozyme, Benzonase [®]	37900
Qprotome Mammalian Protein Prep Kit	For approximately 100 protein preparations from cultured mammalian cells: Buffer, Reagents, Protease Inhibitor Solution, Benzonase [®]	37901
Qproteome Total Glycoprotein Kit	For 6 total glycoprotein preps: Buffers, Lectin Spin Columns (6), Detergent Solution, Protease Inhibitor Solution, Collection Tubes (6 x 2 ml)	37541
Qproteome Mannose Glycoprotein Kit	For 6 mannose glycoprotein preps: ConA, GNA, and LCH Lectin Spin Columns (2 each); Buffers; Detergent Solution; Protease Inhibitor Solution; Collection Tubes (6 x 2 ml)	37551

Product	Contents	Cat. no.
Qproteome Sialic Glycoprotein Kit	For 6 sialic acid glycoprotein preps: WGA, SNA, and MAL Lectin Spin Columns (2 each); Buffers; Detergent Solution; Protease Inhibitor Solution; Collection Tubes (6 x 2 ml)	37561
Qproteome O-Glycan Glycoprotein Kit	For 6 O-glycan glycoprotein preps: ALL and PNA Lectin Spin Columns (3 each); Buffers; Protease Inhibitor Solution; Collection Tubes (6 x 2 ml)	37571
Qproteome Soluble Protein Separation Kit	For 10 soluble protein fractionations: Fractionation Buffer, Precipitation Reagents, Protease Inhibitor Solution, Benzonase [®]	37512
Qproteome Cell Compartment Kit	For 10 subcellular fractionations: Extraction buffers, Protease Inhibitor Solution, Benzonase [®]	37502

QIAGEN Companies

Please see the back cover for contact information for your local QIAGEN office.

QIAGEN Distributors

Argentina

Tecnolab S.A.
Tel: (011) 4555 0010
Fax: (011) 4553 3331
E-mail: info@tecnolab.com.ar
Web site: www.tecnolab.com.ar

Brazil

Uniscience do Brasil
Tel: 011 3622 2320
Fax: 011 3622 2323
E-mail: info@uniscience.com
Web site: www.uniscience.com

Chile

Biosonda SA
Tel: 562 209 6770
Fax: 562 274 5462
E-mail: ventas@biosonda.cl
Web site: www.biosonda.cl

Croatia

INEL Medicinska Tehnika d.o.o.
Tel: (01) 2984-898
Fax: (01) 6520-966
E-mail: inel-medicinska-tehnika@zg.htnet.hr

Cyprus

Scientronics Ltd
Tel: 02-357 22 765416
Fax: 02-357 22 764614
E-mail: a.sarpetsas@biotronics.com.cy

Czech Republic

BIO-CONSULT spol. s.r.o.
Tel/Fax: (420) 2 417 29 792
E-mail: info@bioconsult.cz
Web site: www.bioconsult.cz

Egypt

Clinilab
Tel: 52 57 212
Fax: 52 57 210
E-mail: Clinilab@link.net

Estonia

PEAI-Est OÜ
Tel: (051) 65 830
Fax: (07) 383 360
E-mail: langel@ut.ee

Greece

BioAnalytica S.A.
Tel: (210)-640 03 18
Fax: (210)-646 27 48
E-mail: bioanalyt@hol.gr
Web site: www.bioanalytica.gr

Hungary

Kasztel-Med Co. Ltd.
Tel: (01) 385 3887
Fax: (01) 381 0695
E-mail: info@kasztel.hu
Web site: www.kasztel.hu

India

Genetix
Tel: (011)-2542 1714
or (011)-2515 9346
Fax: (011)-2546 7637
E-mail: genetix@nda.vsnl.net.in

Israel

Westburg (Israel) Ltd.
Tel: 08-6900655
or 1-800 20 22 20 (toll free)
Fax: 08-6900650
E-mail: info@westburg.co.il
Web site: www.westburg.co.il

Korea

LRS Laboratories, Inc.
Tel: (02) 924-86 97
Fax: (02) 924-86 96
E-mail: webmaster@lrslab.co.kr
Web site: www.lrslab.co.kr

Malaysia

RESEARCH BIOLABS SDN. BHD.
Tel: (603)-8070 3101
Fax: (603)-8070 5101
E-mail: biolabs@tm.net.my
Web site: www.researchbiolabs.com

Mexico

Quimica Valaner S.A. de C.V.
Tel: (55) 55 25 57 25
Fax: (55) 55 25 56 25
E-mail: ventas@valaner.com
Web site: www.valaner.com

New Zealand

Biolab Ltd
Tel: (09) 980 6700
or 0800 933 966
Fax: (09) 980 6788
E-mail: biosciences@nzl.biolabgroup.com
Web site: www.biolabgroup.com/nzl

Poland

Syngen Biotech Sp.z.o.o.
Tel: (071) 798 58 50 - 52
Fax: (071) 798 58 53
E-mail: info@syngen.pl
Web site: www.syngen.pl

Portugal

IZASA PORTUGAL, LDA
Tel: (21) 424 7312
Fax: (21) 417 2674
E-mail: consultasbiotec@izasa.es

Saudi Arabia

Abdulla Fouad Co. Ltd.
Tel: (03) 8324400
Fax: (03) 8346174
E-mail: sadiq.omar@abdulla-fouad.com

Singapore

Research Biolabs Pte Ltd
Tel: 6777 5366
Fax: 6778 5177
E-mail: sales@researchbiolabs.com
Web site: www.researchbiolabs.com

Slovak Republic

BIO-CONSULT Slovakia spol. s.r.o.
Tel/Fax: (02) 5022 1336
E-mail: bio-cons@cdicon.sk
Web site: www.bioconsult.cz

Slovenia

MEDILINE d.o.o.
Tel: (01) 830-80-40
Fax: (01) 830-80-70
or (01) 830-80-63
E-mail: mediline@siol.net

South Africa

Southern Cross Biotechnology (Pty) Ltd
Tel: (021) 671 5166
Fax: (021) 671 7734
E-mail: info@scb.co.za

Spain

IZASA, S.A.
Tel: (93) 902.20.30.90
Fax: (93) 902.22.33.66
E-mail: consultasbiotec@izasa.es

Taiwan

TAIGEN Bioscience Corporation
Tel: (02) 2880 2913
Fax: (02) 2880 2916
E-mail: taigen@ms10.hinet.net

Thailand

Theera Trading Co. Ltd.
Tel: (02) 412-5672
Fax: (02) 412-3244
E-mail: theetrad@samart.co.th

Turkey

Medek Medikal Ürünler ve Sağlık Hizmetleri A. S.
Tel: (216) 302 15 80
Fax: (216) 302 15 88
E-mail: makalp@med-ek.com

All other countries

QIAGEN GmbH, Germany

Australia ■ **QIAGEN Pty Ltd** ■ PO Box 641 ■ DONCASTER VIC 3108

Orders 03-9840-9800 ■ Fax 03-9840-9888 ■ Technical 1-800-243-066

Austria ■ **QIAGEN Vertriebs GmbH** ■ Löwengasse 47/Top 6 ■ A-1030 Wien

Orders 0800/28-10-10 ■ Fax 0800/28-10-19 ■ Technical 0800/28-10-11

Belgium ■ **QIAGEN Benelux B.V.** ■ Spoorstraat 50 ■ 5911 KJ Venlo ■ The Netherlands

Orders 0800-79612 ■ Fax 0800-79611 ■ Technical 0800-79556

Canada ■ **QIAGEN Inc.** ■ 2800 Argentia Road ■ Unit 7 ■ Mississauga ■ Ontario ■ L5N 8L2

Orders 800-572-9613 ■ Fax 800-713-5951 ■ Technical 800-DNA-PREP (800-362-7737)

China ■ **QIAGEN China Representative Office** ■ Room 2213 ■ POS PLAZA ■ Century Avenue 1600 ■

Pudong ■ Shanghai 200122 ■ Telephone 021-51162555 ■ Fax 021-51162500

France ■ **QIAGEN S.A.** ■ 3 avenue du Canada ■ LP 809 ■ 91974 COURTABOEUF CEDEX

Orders 01-60-920-920 ■ Fax 01-60-920-925 ■ Technical 01-60-920-930

Germany ■ **QIAGEN GmbH** ■ QIAGEN Strasse 1 ■ 40724 Hilden

Orders 02103-29-12000 ■ Fax 02103-29-22000 ■ Technical 02103-29-12400

Italy ■ **QIAGEN S.p.A.** ■ Via Grosio, 10/10 ■ 20151 Milano

Orders 02-33430411 ■ Fax 02-33430426 ■ Technical 02-33430414

Japan ■ **QIAGEN K.K.** ■ Forefront Tower II ■ 13-1, Kachidoki 3 Chome ■ Chuo-ku, Tokyo 104-0054

Telephone 03-5547-0811 ■ Fax 03-5547-0818 ■ Technical 03-5547-0811

Luxembourg ■ **QIAGEN Benelux B.V.** ■ Spoorstraat 50 ■ 5911 KJ Venlo ■ The Netherlands

Orders 8002-2076 ■ Fax 8002-2073 ■ Technical 8002-2067

The Netherlands ■ **QIAGEN Benelux B.V.** ■ Spoorstraat 50 ■ 5911 KJ Venlo

Orders 0800-0229592 ■ Fax 0800-0229593 ■ Technical 0800-0229602

Switzerland ■ **QIAGEN AG** ■ Garstligweg 8 ■ 8634 Hombrechtikon

Orders 055-254-22-11 ■ Fax 055-254-22-13 ■ Technical 055-254-22-12

UK and Ireland ■ **QIAGEN Ltd.** ■ QIAGEN House ■ Fleming Way ■ Crawley ■ West Sussex, RH10 9NQ

Orders 01293-422-911 ■ Fax 01293-422-922 ■ Technical 01293-422-999

USA ■ **QIAGEN Inc.** ■ 27220 Turnberry Lane ■ Valencia ■ CA 91355

Orders 800-426-8157 ■ Fax 800-718-2056 ■ Technical 800-DNA-PREP (800-362-7737)

