

**Duotech** srl "quo fata vocant"

Via Monte Spluga 31 - 20021 Baranzate (Milano)

[www.duotech.it](http://www.duotech.it) [info@duotech.it](mailto:info@duotech.it)

Tel. +39.0233106630 r.a. fax +39.0233106640



# Mobicols - Plastic Columns

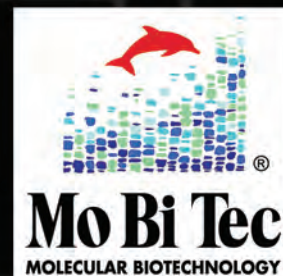
## Tools for diverse Lab Applications

SEPARATION

ISOLATION

TREATMENT

PURIFICATION





- Chapter I**    **Mobicol – A Practical Tool for Every Lab**
- Chapter II**    **Spin Columns for your own Matrices or other Compounds**
- Chapter III**    **Matrix-filled MobiSpin Columns for Purification of Nucleic Acids**
- Chapter IV**    **Mobicols with Enzymatically Active Matrices**

Mobicols are versatile columns which can be used for many lab tasks. They are compatible with sample sizes from very small to very large volumes, and can handle small or large wash buffer volumes **(Chapter I)**.

Mobicols can be used to make your own spin/affinity columns with the matrix of your choice **(Chapter II)**.

Used as a Mini Column they feature Luer-lock connections on top and bottom. Using the Luer-lock adapter, a syringe can be connected to the top of the Mobicol as a reservoir for sample or buffer application. To enable a connection to larger reservoirs, a Luer-adapter is available to attach tubing to the top cap. As Spin Columns, Mobicols are placed in a 1.5 ml or 2 ml microtube and can be centrifuged in a microcentrifuge.

Spin Columns can be used for the purification of biomolecules from small contaminants (e.g. desalting) **(Chapter III)**.

Applying Mini Columns, you can take advantage of highly active immobilized enzymes, packed into small Compact Reaction Columns (CRC) **(Chapter IV)**. With more than 20 years of experience in immobilizing native proteins and active enzymes, MoBiTec offers a custom immobilization service at very attractive conditions.



# Chapter I

**Mobicol – A Practical Tool for Every Lab**





# Mobicol "F"

Mobicols are 700 µl columns with particular handling advantages:

- Mobicols are compatible with laboratory standards
- Mobicols can be centrifuged in microcentrifuges
- Mobicols have a Luer-lock adapter
- Smallest und largest volumes can be treated easily



Luer-lock cap

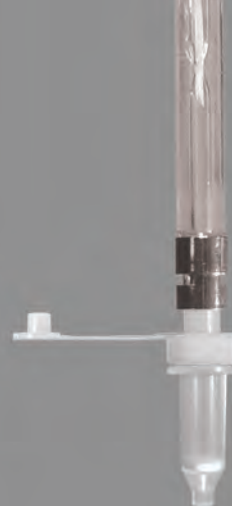
closing cap

35 µm filter

plug



Syringe as reservoir connected to Luer-lock cap of Mobicol



Adapter M3001 tubing to Mobicol

Filter of 35 µm pore size is inserted. A separately available upper filter (Order No. M523515) conveniently caps the resin bed.

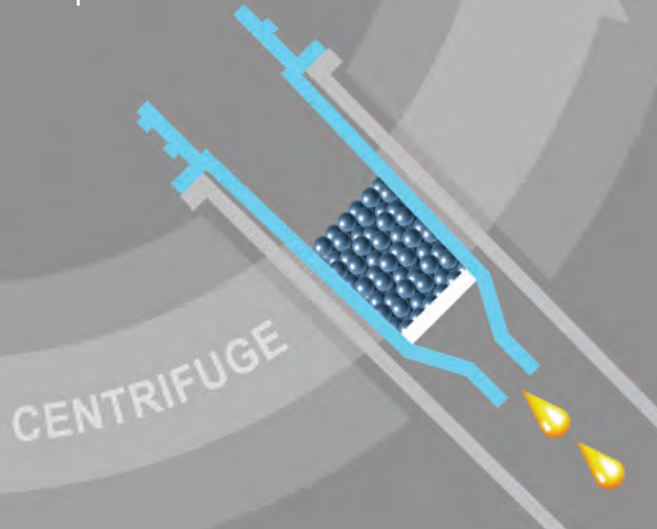
Mobicol "F" can be closed tightly with a screw-on cap and a snap-off plug at the outlet.

Mobicol "F" can be autoclaved at 110 °C for 10 minutes.

**Mobicol "F"**  
with 35 µm filter in  
2 ml collection tube



**Mobicol "F"**  
can also be used  
as Spin Column





## Mobicol "F"

M105035F



Mobicol "F" M105035F with fixed snap-off plug for the outlet is supplied with 2 caps: a Luer-lock cap and a screw cap. The Mobicol "F" can be closed tightly with screw cap and plug. A filter of 35 µm pore size is inserted; an additional upper filter (35 µm) is available.

### Mobicol "F" is approved

- Hundreds of satisfied customers are using Mobicols already for many years
- Many of referenced applications

### Mobicol "F" is versatile

- Can be used in flow-through mode or as Spin Column
- Useful for size exclusion or affinity purification of compounds like nucleic acids and proteins (native or tagged)
- Compatible with small and large sample volumes
- Upper filters of 35 µm pore size are separately available
- Screw cap with rubber seal
- Mobicol can be filled and stored or shipped without leakage
- Reverse side of snap-off plug can be used for resealing

### Mobicol "F" is compatible with lab standards

- Can be centrifuged in standard microcentrifuge
- Luer adapters are available for inlet and outlet tubing
- Luer-lock cap is compatible to Luer syringes
- Autoclavable with inserted filter at 110 °C for 10 min

## Applications

Use it with affinity matrix for:

- **Purification of tagged proteins**
- **Purification of peptides**
- **Purification of native proteins**  
using immobilized IgG, NTP-binding proteins (e.g. kinases), amino acid binding proteins,  $\beta$ -lactamase, Igase, peroxidase, EPO (erythropoietin),  $\beta$ -galactosidase
- **Fractionation and purification of antibodies**  
using protein A/G, immobilized epitopes, peptide matrix
- **Purification of glycoproteins and carbohydrates**
- **Purification of fatty acids and fatty acid binding proteins**
- **Whole mount embryo *in situ* hybridization**

### Order information

Order No.	Description	Amount
<b>M105035F</b>	Mobicol "F" with fixed outlet plug, inserted 35 µm filter and 2 different screw caps	50
<b>M1050</b>	Mobicol "F" with fixed outlet plug and 2 different screw caps, without filters	50
<b>M1053</b>	Mobicol "F" with fixed outlet plug and screw cap, without filter	50
<b>M2210</b>	Filter (large) 10 µm pore size, for Mobicol M1002, M1003, M1050 & M1053	50
<b>M523515</b>	Filter (large) 35 µm pore size, for Mobicol M1002, M1003, M1050 & M1053	50
<b>M2190</b>	Filter (small) 90 µm pore size, for Mobicol M1002, M1003, M1050 & M1053	50

For column accessories see page 7.

Also available prefilled with immobilized enzymes:

Immobilized proteases, nucleases, alk. phosphatase,  $\beta$ -galactosidase, or a compound of your choice on request (see chapter IV).

**Mobicol "Classic"**

Use for Affinity Chromatography (M1002)



with inserted filters (large)

**Mobicol "Classic"**

Use as Spin Column (M1003)



with inserted filters (small)

The Mobicol "Classic" is supplied with two screw caps (Luer-lock cap and screw cap) and a plug for the outlet. Small and large filters with different diameters (10  $\mu$ m, 35  $\mu$ m and 90  $\mu$ m) are separately available, so you can use the filter of your choice. With this features the Mobicol "Classic" can be used for a variety of applications.

**Filters and insertion tools**

(small)

(large)

**Mobicol "Classic"**

Order No.	Description	Amount
<b>M1002</b>	Mobicol "Classic" with 2 different screw caps, without filters	50
<b>M1003</b>	Mobicol "Classic" with 1 screw cap, without filters	50
<b>M2110</b>	Filter (small) 10 $\mu$ m pore size, for Mobicol M1002, M1003, M1050 & M1053	50
<b>M513515</b>	Filter (small) 35 $\mu$ m pore size, for Mobicol M1002, M1003, M1050 & M1053	50
<b>M2190</b>	Filter (small) 90 $\mu$ m pore size, for Mobicol M1002, M1003, M1050 & M1053	50
<b>M2210</b>	Filter (large) 10 $\mu$ m pore size, for Mobicol M1002, M1003, M1050 & M1053	50
<b>M523515</b>	Filter (large) 35 $\mu$ m pore size, for Mobicol M1002, M1003, M1050 & M1053	50
<b>M2290</b>	Filter (large) 90 $\mu$ m pore size, for Mobicol M1002, M1003, M1050 & M1053	50

## Accessories for Affinity Chromatography Columns

**M3003**  
Luer adapter (M)  
for the top-cap



**M3004**  
Luer adapter (F) for outlet



**M3006**  
Stopcock, one-way,  
Luer-lock (F)



**M3007**  
Luer-lock adapter (M/M)



**M3001**  
Tubing-to-Luer-lock  
adapter

connected with

**M1002**  
Mobicol „Classic“



Large filter inserted  
(not enclosed)

The polypropylene Luer adapter (M) (M3003) for tubing connection to the inlet of the column allows a flow-through mode at low pressure, whereas the Luer adapter (F) (M3004) is suitable for a tubing connection to the outlet of the column. Furthermore the Luer adapter (M/M) (M3007) facilitates the connection of the inlet to a Luer-lock system.

The Tubing-to-Luer-lock adapter (M3001) enables a safe connection from a pump to the inlet of a Mobicol or a 2.5 ml column.

The one-way stopcock, Luer-lock (F) (M3006) is a convenient tool for controlling flow-through at the in- or outlet.

### Column Accessories

Order No.	Description	Amount
<b>M3001</b>	Tubing-to-Luer-lock adapter (material outside: metal; inside: glass and Teflon; with 1 m Teflon tubing) for Mobicols or laboratory columns	1
<b>M3002</b>	Luer adapter, (M+F) 10 each	20
<b>M3003</b>	Luer adapter for the top-cap (M)	20
<b>M3004</b>	Luer adapter for outlet (F)	20
<b>M3007</b>	Luer-lock adapter (M/M)	1
<b>M3005</b>	Bottom plugs for Mobicols	50
<b>M3006</b>	Stopcocks, one-way, Luer-lock (F)	4
<b>M3009</b>	Luer-lock caps	50



## References

Jinhi Ahn, Jason T. Wong, and Robert S. Molday The Effect of Lipid Environment and Retinoids on the ATPase Activity of ABCR, the Photoreceptor ABC Transporter Responsible for Stargardt Macular Dystrophy *J. Biol. Chem.*, Jun 2000; 275: 20399.

Véronique Amiard, Annette Morvan-Bertrand, Jean-Pierre Billard, Claude Huault, Felix Keller, and Marie-Pascale Prud'homme Fructans, But Not the Sucrosyl-Galactosides, Raffinose and Loliose, Are Affected by Drought Stress in Perennial Ryegrass Plant Physiology, Aug 2003; 132: 2218.

Pierre-Olivier Angrand, Inmaculada Segura, Pamela Völkel, Sonja Ghidelli, Rebecca Terry, Miro Brajenovic, Kristina Vintersten, Rüdiger Klein, Giulio Superti-Furga, Gerard Drewes, Bernhard Kuster, Tewis Bouwmeester, and Amparo Acker-Palmer Transgenic Mouse Proteomics Identifies New 14-3-3-associated Proteins Involved in Cytoskeletal Rearrangements and Cell Signaling *Mol. Cell. Proteomics*, Dec 2006; 5: 2211 - 2227.

Denis Bayle, Sabine Wängler, Thomas Weitzenegger, Wolfram Steinhilber, Jürgen Volz, Michael Przybylski, Klaus P. Schäfer, George Sachs, and Klaus Melchers Properties of the P-Type ATPases Encoded by the copAP Operons of *Helicobacter pylori* and *Helicobacter felis* *J. Bacteriol.*, Jan 1998; 180: 317 - 329.

Miro Brajenovic, Gerard Joberty, Bernhard Küster, Tewis Bouwmeester, and Gerard Drewes Comprehensive Proteomic Analysis of Human Par Protein Complexes Reveals an Interconnected Protein Network *J. Biol. Chem.*, Mar 2004; 279: 12804 - 12811.

Jan Brix, Klaus Dietmeier, and Nikolaus Pfanner Differential Recognition of Preproteins by the Purified Cytosolic Domains of the Mitochondrial Import Receptors Tom20, Tom22, and Tom70 *J. Biol. Chem.*, Aug 1997; 272: 20730.

Jan Brix, Stefan Rüdiger, Bernd Bukau, Jens Schneider-Mergener, and Nikolaus Pfanner Distribution of Binding Sequences for the Mitochondrial Import Receptors Tom20, Tom22, and Tom70 in a Presequence-carrying Preprotein and a Non-cleavable Preprotein *J. Biol. Chem.*, Jun 1999; 274: 16522.

Alejandro Ferrando, Zsuzsana Koncz-Kálmán, Rosa Farràs, Antonio Tiburcio, Jeff Schell, and Csaba Koncz Detection of in vivo protein interactions between Snf1-related kinase subunits with intron-tagged epitope-labelling in plants cells *Nucleic Acids Res.*, Sep 2001; 29: 3685 - 3693.

J. Fleck Chemical Fate of a Metamorphic Inducer in Larvae-like Buds of the Cnidarian *Cassiopea andromeda* *Biol. Bull.*, Feb 1998; 194: 83.

Albrecht Gruhler, Jesper V. Olsen, Shabaz Mohammed, Peter Mortensen, Nils J. Færgeman, Matthias Mann, and Ole N. Jensen Quantitative Phosphoproteomics Applied to the Yeast Pheromone Signaling Pathway *Mol. Cell. Proteomics*, Mar 2005; 4: 310 - 327.

Klaus Hellmuth, Denise M. Lau, F. Ralf Bischoff, Markus Künzler, Ed Hurt, and George Simos CELL AND ORGANELLE STRUCTURE AND ASSEMBLY: Yeast Los1p Has Properties of an Exportin-Like Nucleocytoplasmic Transport Factor for tRNA .

Munir Iqbal, Emma Poole, Stephen Goodbourn, and John W. McCauley Role for Bovine Viral Diarrhea Virus Erns Glycoprotein in the Control of Activation of Beta Interferon by Double-Stranded RNA *J. Virol.*, Jan 2004; 78: 136 - 145.

Josef Wissing, Lothar Jänsch, Manfred Nimtz, Guido Dieterich, Renate Hornberger, György Kéri, Jürgen Wehland, and Henrik Daub Proteomics Analysis of Protein Kinases by Target Class-selective Prefractionation and Tandem Mass Spectrometry *Mol. Cell. Proteomics*, Mar 2007; 6: 537 - 547.

Irvine SQ Whole-mount in situ hybridization of small invertebrate embryos using laboratory mini-columns *Biotechniques*, Dec 2007; 43(6): 764, 766,768.

Florian Krötz, Barbara Engelbrecht, Martin A. Buerkle, Florian Bassermann, Hanna Bridell, Torsten Gloe, Justus Duyster, Ulrich Pohl, and Hae-Young Sohn The Tyrosine Phosphatase, SHP-1, Is a Negative Regulator of Endothelial Superoxide Formation *J. Am. Coll. Cardiol.*, May 2005; 45: 1700 - 1706.

Jelle Van Leene, Hilde Stals, Dominique Eeckhout, Geert Persiau, Eveline Van De Slijke, Gert, Van Isterdael, Annelies De Clercq, Eric Bonnet, Kris Laukens, Noor Remmerie, Kim Hendrickx, Thomas De Vijlder, Azmi Abdelkrim, Anne Pharazyn, Harry Van Onckelen, Dirk Inzé, Erwin Witters, and Geert De Jaeger A Tandem Affinity Purification-based Technology Platform to Study the Cell Cycle Interactome in *Arabidopsis thaliana* *Mol. Cell. Proteomics*, Jul 2007; 6: 1226 - 1238.

Jérémy Lothier, Bertrand Lasseur, Katrien Le Roy, André Van Laere, Marie-Pascale Prud'homme, Philippe Barre, Wim Van den Ende, and Annette Morvan-Bertrand Cloning, gene mapping, and functional analysis of a fructan 1-exohydrolase (1-FEH) from *Lolium perenne* implicated in fructan synthesis rather than in fructan mobilization *J. Exp. Bot.*, Jun 2007; 58: 1969 - 1983.

A. Magrelli, K Langenkemper, C Dehio, J Schell, and A Spena Splicing of the rolA transcript of *Agrobacterium rhizogenes* in *Arabidopsis* *Science*, Dec 1994; 266: 1986 - 1988.

Heidmarie Müllner, Günter Deutsch, Erich Leitner, Elisabeth Ingolic, and Günther Daum YEH2/YLR020c Encodes a Novel Steryl Ester Hydrolase of the Yeast *Saccharomyces cerevisiae* *J. Biol. Chem.*, Apr 2005; 280: 13321 - 13328.

Karsten Niederhoff, Nadja M. Meindl-Beinker, Daniela Kerksen, Uta Perband, Antje Schäfer, Wolfgang Schliebs, and Wolf-H. Kunau Yeast Pex14p Possesses Two Functionally Distinct Pex5p and One Pex7p Binding Sites *J. Biol. Chem.*, Oct 2005; 280: 35571 - 35578.

Stephenie Paine-Saunders, Beth L. Viviano, Aris N. Economides, and Scott Saunders Heparan Sulfate Proteoglycans Retain Noggin at the Cell Surface. A POTENTIAL MECHANISM FOR SHAPING BONE MORPHOGENETIC PROTEIN GRADIENTS *J. Biol. Chem.*, Jan 2002; 277: 2089 - 2096.

K. Pfeifer, M Bachmann, HC Schroder, BE Weiler, D Ugarkovic, T Okamoto, and WE Muller Formation of a small ribonucleoprotein particle between Tat protein and trans-acting response element in human immunodeficiency virus-infected cells *J. Biol. Chem.*, Aug 1991; 266: 14620 - 14626.

Werner G. Purschke, Falko Radtke, Frank Kleinjung, and Sven Klussmann A DNA Spiegelmer to staphylococcal enterotoxin B *Nucleic Acids Res.*, Jun 2003; 31: 3027 - 3032.

Juri Rappsilber, Paul Ajuh, Angus I. Lamond, and Matthias Mann SPF30 Is an Essential Human Splicing Factor Required for Assembly of the U4/U5/U6 Tri-small Nuclear Ribonucleoprotein into the Spliceosome *J. Biol. Chem.*, Aug 2001; 276: 31142 - 31150.

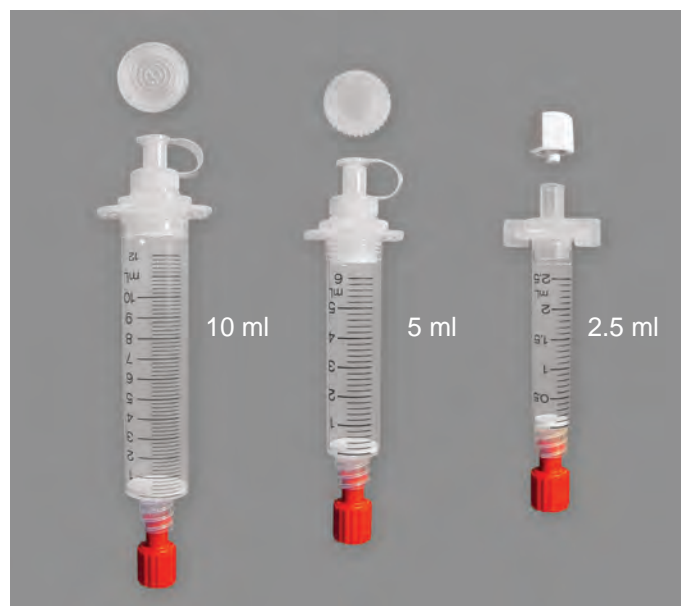
B. Ashok Reddy, Prashanth Kumar Bajpe, Andrew Bassett, Yuri M. Moshkin, Elena Kozhevnikova, Karel Bezstarosti, Jeroen A.A. Demmers, Andrew A. Travers, and C. Peter Verrijzer Drosophila Transcription Factor Tramtrack69 Binds MEP1 to Recruit the Chromatin Remodeler NuRD *Mol. Cell. Biol.*, Aug 2010; 30: 1128/ MCB.00266-10.

Ulrich Rothbauer, Kourosh Zolghadr, Serge Muyldermans, Aloys Schepers, M. Cristina Cardoso, and Heinrich Leonhardt A Versatile Nanotrap for Biochemical and Functional Studies with Fluorescent Fusion Proteins *Mol. Cell. Proteomics*, Feb 2008; 7: 282 - 289.



## Related Products

For volumes larger than 1 ml (Mobicols), 2.5 ml, 5 ml and 10 ml columns are available with filters inserted. These syringe based columns have Luer-lock adapters at their in- and outlet so that they have the same handling advantages as Mobicols.



## Laboratory columns

Order No.	Description	Amount
S10121	2.5 ml columns (10 µm filter pore size)	20
S1012	2.5 ml columns (35 µm filter pore size)	20
S10129	2.5 ml columns (90 µm filter pore size)	20
S10131	5 ml columns (10 µm filter pore size)	20
S1013	5 ml columns (35 µm filter pore size)	20
S10139	5 ml columns (90 µm filter pore size)	20
S10141	10 ml columns (10 µm filter pore size)	20
S1014	10 ml columns (35 µm filter pore size)	20
S10149	10 ml columns (90 µm filter pore size)	20
S1210	Upper filter 10 µm for 2.5 ml columns	20
S1235	Upper filter 35 µm (thickness 1.5 mm) for 2.5 ml columns	20
S123532	Upper filter 35 µm (thickness 3.2 mm) for 2.5 ml columns	20
S1290	Upper filter 90 µm for 2.5 ml columns	20
S1310	Upper filter 10 µm for 5 ml columns	20
S1335	Upper filter 35 µm (thickness 1.5 mm) for 5 ml columns	20
S133532	Upper filter 35 µm (thickness 3.2 mm) for 5 ml columns	20
S1390	Upper filter 90 µm for 5 ml columns	20
S1410	Upper filter 10 µm for 10 ml columns	20
S143532	Upper filter 35 µm (thickness 3.2 mm) for 10 ml columns	20
S1490	Upper filter 90 µm for 10 ml columns	20

## Column sets

Order No.	Description	Amount
S10011	Set: Mobicol "F" M1050; 2.5 ml ; 5 ml and 10 ml columns (10 µm pore size filter)	3 each
S1001	Set: Mobicol "F" M1050; 2.5 ml ; 5 ml and 10 ml columns (35 µm pore size filter)	3 each
S10019	Set: Mobicol "F" M1050; 2.5 ml ; 5 ml and 10 ml columns (90 µm pore size filter)	3 each
S10031	Set upper filters 10 µm pore size for S10011	1 set
S1003	Set upper filters 35 µm pore size for S1001	1 set
S10039	Set upper filters 90 µm pore size for S10019	1 set



## Chapter II

**Spin Columns for your own Matrices  
or other Compounds**

## Advantages

**Spin column chromatography offers many advantages over traditional liquid chromatography:**

- Comes empty - fill in your own material (matrix, gel slice etc.)
- Easy handling: load sample, spin and collect the purified product
- No sample dilution
- Reproducible results with fast protocols
- One application in less than 4 minutes
- Numerous samples can be processed simultaneously
- Large number of applications
- Compatible with laboratory standard

## Background

The MobiSpin column "F" is designed for a wide variety of applications for nucleic acid and protein extraction and purification. When choosing the MobiSpin column "F" for a particular application, a suitable resin must be selected and filled into the mini-column. For an appropriate resin selection, impurity versus sample size and the anticipated yield must be considered.

The MobiSpin column "F" comes with inserted small filter of 10 µm pore size, screw cap and snap-off plug. It can be used without matrix (e.g. to extract trypsinated proteins out of a gel slice for Mass Spec analysis) or filled with a matrix of your choice (see also Chapter III).

**Gel matrix or gel slice or other compound**

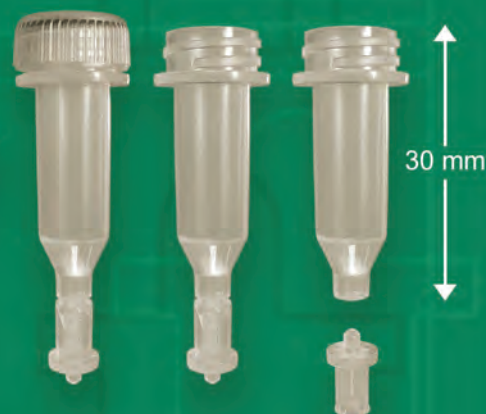
**Fill in your own material**

## MobiSpin Column "F"

can be centrifuged  
in 1.5 ml or 2 ml tubes

centrifuge

Illustration enlarged - fits all common centrifuges

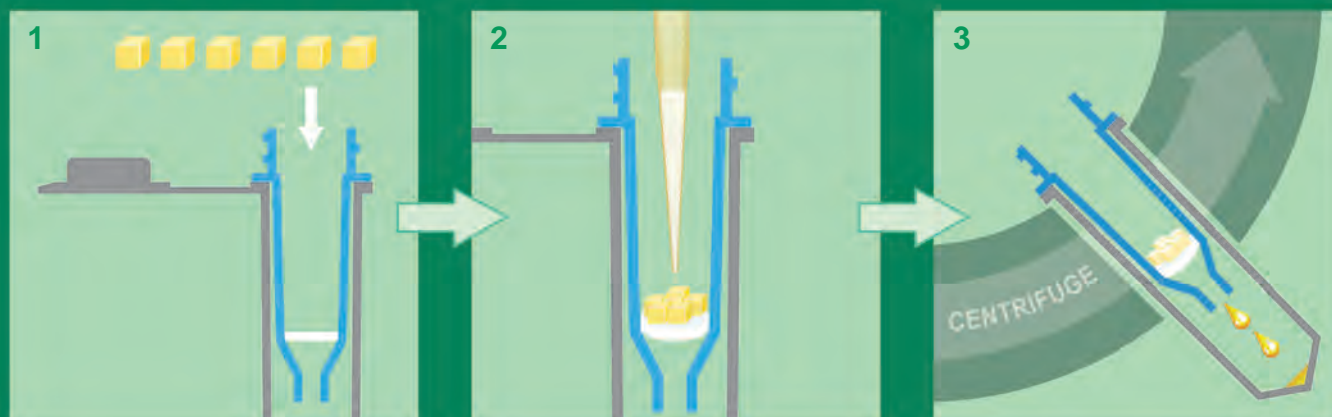


The MobiSpin column is autoclavable at 110 °C for 10 min.  
Close the column with the tightly closing screw cap.  
Snap off the outlet plug and use its reverse side for closing.



## Special Application

## Preparation of trypsinated protein fragments for Mass Spec



For mass spectrometry analysis, proteins are separated by polyacrylamide gel electrophoresis (PAGE) and a single gel slice containing a protein of interest is cut out of the gel. The slice is cut into pieces of approximately 2 x 2 mm and then placed onto the inserted filter of a MobiSpin column "F" (M105210S) (1). After several washing and drying steps involving centrifugation, trypsin is added to the gel pieces inside of the column, diffuses into them and digests the protein overnight at 37 °C into small fragments (2). The protein fragments are eluted by a few spinning steps (3). After drying the collected spin elutes can be analyzed by mass spectrometry. The complete protocol is available on our web page ([www.mobitec.com](http://www.mobitec.com)). The peptide mass fingerprints provided by this protocol are consistent with results obtained by other common mass spec sample preparations.

This protocol will be suitable for proteases other than trypsin as well.

## MobiSpin Column "F"

Order No.	Description	Amount
M105010S	MobiSpin Column "F" with fixed outlet plug, inserted small 10 µm filter and screw cap	50
M105210S	MobiSpin Column "F" with fixed outlet plug, inserted large 10 µm filter and screw cap	50



## References

Demian Cazalla, Kathryn Newton, and Javier F. Cáceres A Novel SR-Related Protein Is Required for the Second Step of Pre-mRNA Splicing *Mol. Cell. Biol.*, Apr 2005; 25: 2969 - 2980.

B. K. Choi, B J Paster, F E Dewhirst, and U. B. Göbel Diversity of cultivable and uncultivable oral spirochetes from a patient with severe destructive periodontitis. *Infect. Immun.*, May 1994; 62: 1889 - 1895.

Holger Eubel, Etienne H. Meyer, Nicolas L. Taylor, John D. Bussell, Nicholas O'Toole, Joshua L. Heazlewood, Ian Castleden, Ian D. Small, Steven M. Smith, and A. Harvey Millar Novel Proteins, Putative Membrane Transporters, and an Integrated Metabolic Network Are Revealed by Quantitative Proteomic Analysis of Arabidopsis Cell Culture Peroxisomes *Plant Physiology*, Dec 2008; 148: 1809 - 1829.

J. Hedegaard, SA Steffensen, N Norskov-Lauritsen, KK Mortensen, and HU Sperling-Petersen Identification of Enterobacteriaceae by partial sequencing of the gene encoding translation initiation factor 2 *Int. J. Syst. Bacteriol.*, Oct 1999; 49: 1531.

Yoko Kita, Yoshiaki Miura, Jun-ichi Furukawa, Mika Nakano, Yasuro Shinohara, Masahiro Ohno, Akio Takimoto, and Shin-Ichiro Nishimura Quantitative Glycomics of Human Whole Serum Glycoproteins Based on the Standardized Protocol for Liberating N-Glycans *Mol. Cell. Proteomics*, Aug 2007; 6: 1437 - 1445.

Angeles Ortega, Martina Niksic, Angela Bachi, Matthias Wilm, Lucas Sánchez, Nicholas Hastie, and Juan Valcárcel Biochemical Function of Female-Lethal (2)D/Wilms' Tumor Suppressor-1-associated Proteins in Alternative Pre-mRNA Splicing, *J. Biol. Chem.*, Jan 2003; 278: 3040 - 3047.

Shaun Peters, Sagadevan G. Mundree, Jennifer A. Thomson, Jill M. Farrant, and Felix Keller Protection mechanisms in the resurrection plant *Xerophyta viscosa* (Baker): both sucrose and raffinose family oligosaccharides (RFOs) accumulate in leaves in response to water deficit *J. Exp. Bot.*, Jun 2007; 58: 1947 - 1956.

Stephanie Pfander, Roberto Fiammengio, Srecko I. Kirin, Nils Metzler-Nolte, and Andres Jäschke Reversible site-specific tagging of enzymatically synthesized RNAs using aldehyde-hydrazine chemistry and protease-cleavable linkers *Nucleic Acids Res.*, Feb 2007; 35: e25.

Sonja Reiland, Gaëlle Messerli, Katja Baerenfaller, Bertran Gerrits, Anne Endler, Jonas Grossmann, Wilhelm Gruissem, and Sacha Baginsky Large-Scale Arabidopsis Phosphoproteome Profiling Reveals Novel Chloroplast Kinase Substrates and Phosphorylation Networks *Plant Physiology*, Jun 2009; 150: 889 - 903.

Symeon Siniosoglou, Ed C. Hurt, and Hugh R. B. Pelham Psr1p/Psr2p, Two Plasma Membrane Phosphatases with an Essential DXDX(T/V) Motif Required for Sodium Stress Response in Yeast *J. Biol. Chem.*, Jun 2000; 275: 19352.

W. Voos, B. D. Gambill, S. Laloraya, D. Ang, E. A. Craig, and N. Pfanner Mitochondrial GrpE is present in a complex with hsp70 and preproteins in transit across membranes. *Mol. Cell. Biol.*, Oct 1994; 14: 6627 - 6634.

F. T. Vreede and G. G. Brownlee Influenza Virion-Derived Viral Ribonucleoproteins Synthesize both mRNA and cRNA *In Vitro J. Virol.*, Mar 2007; 81: 2196 - 2204.

Juan Zou, Lorna Henderson, Vicky Thomas, Patricia Swan, A. Neil Turner, and Richard. G. Phelps Presentation of the Goodpasture Autoantigen Requires Proteolytic Unlocking Steps That Destroy Prominent T Cell Epitopes *J. Am. Soc. Nephrol.*, Mar 2007; 18: 771 - 779.



# Chapter III

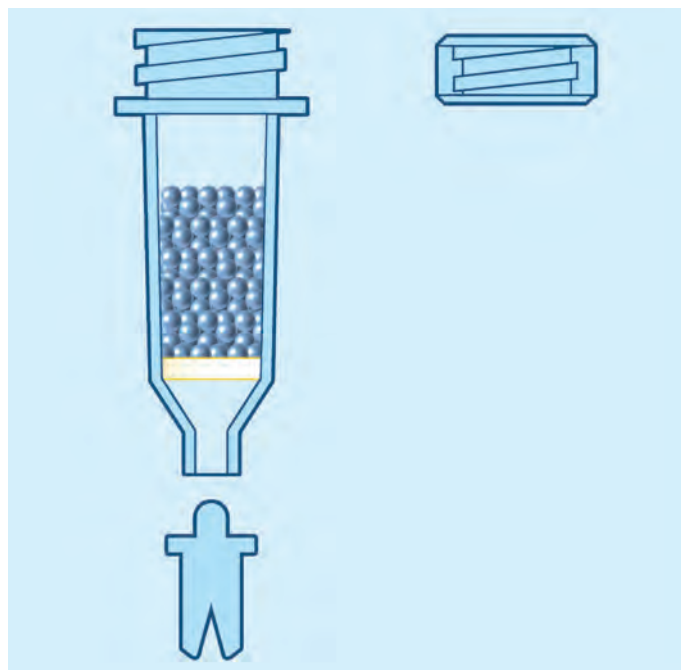
## Matrix-filled MobiSpin Columns for Purification of Nucleic Acids





## Pre-packed MobiSpin Columns for Separation of DNA and RNA from small Contaminants

The MobiSpin Columns are designed for a wide variety of separation tasks. In this system, a gel matrix consisting of porous particles in aqueous solution is packed into a column and then used for the separation of biomolecules. The filled-in matrix pore size determines which molecules are small enough to enter the matrix beads and which molecules are too large and thus remain outside. Molecules which are larger than the matrix pore size and which cannot enter the matrix beads, appear only in the matrix void volume between the beads. From the inter-bead space they can be recovered by spinning the column in a collection tube in a benchtop centrifuge. Strong spinning elutes the column dead volume without washing and thus without dilution, also in small volumes. Smaller biomolecules located inside of the matrix beads are not eluted by spinning.



### Column Usage Guidelines

- 20x rule: For obtaining optimal results, the smallest product being purified should be at least 20 times larger than the largest impurity. A difference in size smaller than 20-fold may affect either purity or yield.
- The larger the pore size of the resin, the greater the purity and the lower the yield of the resulting product. Gel filtration matrices with larger pore size (S-300, S-400) usually retain more product than matrices with smaller pore size (e.g. S-200).
- Non-specific binding: The MobiSpin Columns exhibit only insignificant non-specific binding, allowing purification of samples in the nanogram range.
- Retention: With increasing molecular weight, relative retention decreases.
- For optimizing please use the Sample Volume Guide on page 19.



## Application: Removal of small Molecules

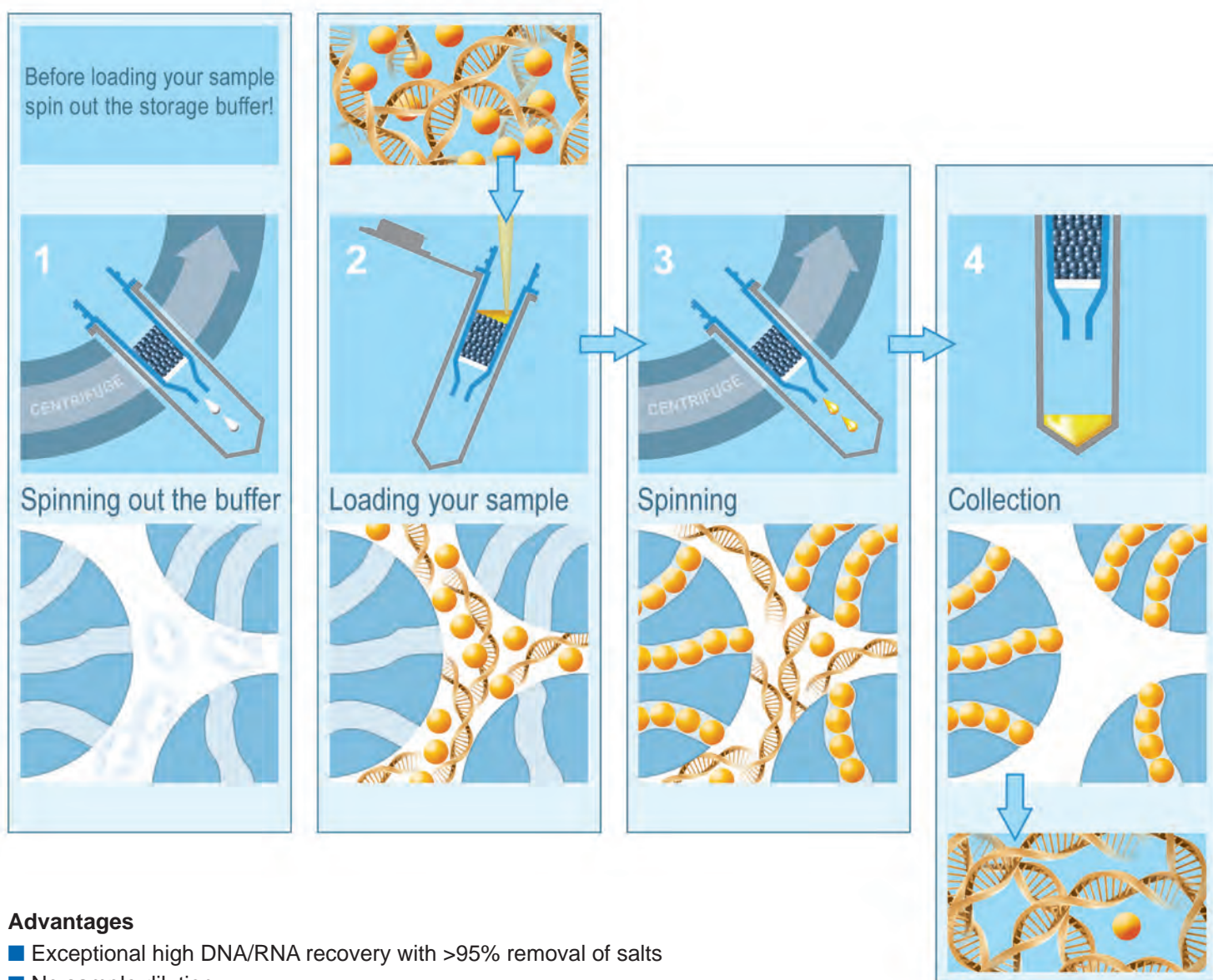
### Desalting, removal of ions

Removing salts and other small molecules from biomolecules is often essential for downstream processes. MobiSpin Columns quickly remove small molecules like salt ions from DNA/RNA. The MobiSpin G-50 columns contain a high-performance size-exclusion matrix for effective salt removal and excellent biomolecule recovery. MobiSpin G-50 matrix with a pore size of 700 Da allows hydrated salt ions to enter the matrix beads while DNA/RNA (and most other biomolecules) stay outside.

 **Removing excess derivatizing agents** from modified DNA/RNAs

 **Removing unreacted dyes** from fluorescence labeling reactions

 **Removing free radiolabel** from labeled DNA and RNA



### Advantages

- Exceptional high DNA/RNA recovery with >95% removal of salts
- No sample dilution
- No long lasting gravity flow
- No chromatography system
- No column preparation
- Easy handling, multiple sample processing in a few minutes



## Applications

### MobiSpin Columns for Spin Size Exclusion Chromatography

MobiSpin columns with S-200, S-300 or S-400 matrix can be used to purify plasmid DNA, cDNA, PCR fragments or oligonucleotides.

- DNA purification prior to sequencing
- Oligonucleotide purification after removal of free nucleotides
- Low cost plasmid purification

### Oligonucleotides Purification after Synthesis

Oligonucleotides can be purified directly from ammonia deprotection solution using MobiSpin Columns: a synthesized 17-mer oligonucleotide (125 µl) was purified directly using a MobiSpin Column S-200 according to standard protocol. For comparison, a 500 µl aliquot of the same deprotection solution was purified on a Sephadex G-25 DNA Grade drip column. The quality of DNA obtained with the MobiSpin Column as analyzed by electrophoresis was equivalent to that obtained with the drip column, however purification using the MobiSpin Column took only one fifth of the time required to purify the sample on the drip column.

### Removal of <sup>32</sup>P-Labeled Nucleotides

MobiSpin Columns can be used to remove large amounts of unincorporated radiolabeled nucleotides. Using the MobiSpin Columns, a 792 bp endlabeled fragment was purified over each of the three MobiSpin Column types. The unincorporated label from 20 µCi reactions was easily removed from the labeled DNA in all cases. In a second experiment, the same 792 bp fragment was used as a template for oligolabeling. Again, the unincorporated labeled nucleotides from 50 µCi reactions were completely and efficiently removed during purification of the reaction products on each type of MobiSpin Column.

### Removal of DNA Nucleotides after PCR

Using our MobiSpin Columns, after PCR reactions, DNA nucleotides can be recovered over a wide range of sample loading levels.

In our control experiment, 1 ng of labeled pUC 18 DNA in 50 µl sample volume was purified over each of the MobiSpin Columns S-200, S-300 and S-400. The complete removal of unincorporated, labeled nucleotides was achieved with excellent recovery of labeled DNA.

In a second test, various amounts of labeled DNA (25 - 250 ng) from an endlabeled reaction were combined with large quantities of unlabeled nucleotides (25 - 250 ng). The mixtures were purified over each type of MobiSpin Column. Under these conditions, regardless the resin type, the presence of relatively large amounts of nucleotides had virtually no effect on the yield of longer labeled DNA or on the ability of the column to remove unincorporated nucleotides from the reaction mixture.

### Use of MobiSpin S-200 Columns in PCR Template Prep for ssDNA Sequencing

MobiSpin S-200 columns were used in two steps for purifying the PCR template for ssDNA sequencing. Following production of a phosphorylated PCR product, the MobiSpin S-200 column exchanged the buffer and removed the unincorporated nucleotides (1st step). After treating the PCR product with lambda exonuclease (which digests the phosphorylated strand only), a buffer exchange and nucleotide removal were carried out again using a MobiSpin S-200 column (2nd step). After this two-step purification, the purified single-stranded DNA could be sequenced directly.

### DNA Purification prior to Sequencing

Plasmid DNA was prepared using a standard alkaline lysis procedure. The miniprep DNA was denatured to the single-stranded form. Then, a volume of 25 µl was purified on each type of MobiSpin Column prior to sequencing. For comparison, CsCl-purified DNA was sequenced in a parallel reaction. All three types of MobiSpin Columns yielded results comparable to the CsCl-purified template DNA.





## MobiSpin Columns for DNA/RNA purification, separation of larger DNA from mono- and multimers

### Sample Volume Guide for S-200, S-300, S-400

<b>Maximum Contaminant Size: 25-mers</b>				
				S-200 (10 µl)
				S-300 (10 µl)
				S-400 (25-50 µl)
<b>Maximum Contaminant Size: 18-mers</b>				
			S-200 (10 µl)	S-200 (10 µl)
	S-300 (25-50 µl)		S-300 (25-50 µl)	S-300 (25-50 µl)
	S-400 (50-100 µl)		S-400 (50-100 µl)	S-400 (50-100 µl)
<b>Maximum Contaminant Size: 8-mers</b>				
	S-200 (25-50 µl)	S-200 (25-50 µl)	S-200 (10-50 µl)	S-200 (10-50 µl)
	S-300 (25-50 µl)	S-300 (25-75 µl)	S-300 (25-75 µl)	S-300 (25-75 µl)
	S-400 (50-100 µl)	S-400 (50-100 µl)	S-400 (50-100 µl)	
<b>Maximum Contaminant Size: NTPs/salts</b>				
	S-200 (25-50 µl)	S-200 (25-50 µl)	S-200 (10-50 µl)	S-200 (10-50 µl)
	S-300 (25-50 µl)	S-300 (25-75 µl)	S-300 (25-75 µl)	S-300 (25-75 µl)
	S-400 (50-100 µl)	S-400 (50-100 µl)	S-400 (50-100 µl)	
<b>Minimum Product Size:</b>				
>20-mers	>50-mers	>200-mers	>500-mers	

The sample volumes in the table have been tested for each of the matrices S-200, S-300 and S-400. The recommended sample volume, as well as the matrix suited best for your particular application, can be found at the intersection of the minimum product size (horizontal) and the estimated maximum contaminant size (vertical).

For example, if you want to separate an 8-mer contaminant (3rd row) from your 100 bp molecule (50mer < 100 bp < 200mer; 2nd column), please go to the third row and se-

cond column of the table and, dependend on your sample volume, select S-200 (if your sample volume is 25-50 µl), S-300 (if your sample volume is 25-75 µl) or S-400 (if your sample volume is 50-100 µl).

This table is intended to be only a general guideline. The final selection of column type depends on your specific application. If you are uncertain about the optimal volume, apply a sample volume of 50 µl.

### Standard Protocol

- Resuspend the resin in the column by vortexing
- Open the bottom plug and loose the cap one fourth turn
- Place the column in a 1.5 ml screw-cap microcentrifuge tube for support.
- Pre-spin the column 1 minute at 800 x g in a microcentrifuge.
- Place the column in a new 1.5 ml tube and slowly apply the sample (10 -100 µl) to the top of the resin, being carefull not to disturb the bed.
- Spin the column 2 minutes at 800 x g. The purified sample is collected in the bottom of the support tube.

MobiSpin Columns are available with three different Sephacryl® resins: S-200, S-300 and S-400. Furthermore, MoBiTec offers MobiSpin Columns filled with Sephadex® G-50 resin. The matrix selection depends on distinct factors, e.g. the sample volume, the size and three-dimensional structure of the product, the g-forces used in the purification process, the resin used as well as the depth of the resin bed.

® Trademarks are registered by Pharmacia.

## Specifications

**MobiSpin S-200** columns: 20 columns, pre-packed with Sephacryl® S-200 HR resin and equilibrated in 10 mM Tris/HCl, 1 mM EDTA buffer at pH 7.6

**MobiSpin S-300** columns: 20 columns, pre-packed with Sephacryl® S-300 HR resin and equilibrated in 10 mM Tris/HCl, 1 mM EDTA buffer at pH 7.6

**MobiSpin S-400** columns: 20 columns, pre-packed with Sephacryl® S-400 HR resin and equilibrated in 10 mM Tris/HCl, 1 mM EDTA buffer at pH 7.6

**MobiSpin columns set:** 30 pre-packed columns, (10 of each of the above resins) equilibrated in 10 mM Tris/HCl, 1mM EDTA buffer at pH 7.6

**MobiSpin G-50** columns: 20 columns, pre-packed with Sephadex® G-50 resin and equilibrated in 10 mM Tris/HCl, 1mM EDTA, 100 mM NaCl buffer at pH 8.0

**Equilibrated  
and  
ready-to-use**

The column caps and their corresponding labels are color-coded for easy identification:

MobiSpin S-200  
columns:  
**Red cap**

MobiSpin S-300  
columns:  
**Green cap**

MobiSpin S-400  
columns:  
**Yellow cap**

MobiSpin G-50  
columns:  
**Neutral cap**

## Order Information, Shipping and Storage

Order No.	Description	Amount
<b>SCO200</b>	MobiSpin S-200 pre-packed with 400 µl matrix	20 columns
<b>SCO210</b>	MobiSpin S-200 pre-packed with 400 µl matrix	100 columns
<b>SCO300</b>	MobiSpin S-300 pre-packed with 400 µl matrix	20 columns
<b>SCO310</b>	MobiSpin S-300 pre-packed with 400 µl matrix	100 columns
<b>SCO400</b>	MobiSpin S-400 pre-packed with 400 µl matrix	20 columns
<b>SCO410</b>	MobiSpin S-400 pre-packed with 400 µl matrix	100 columns
<b>SCO234</b>	MobiSpin S-200, S-300, S-400 pre-packed with 400 µl matrix	3 x 10 columns
<b>SCO500</b>	MobiSpin G-50 pre-packed with 500 µl matrix	20 columns
<b>SCO510</b>	MobiSpin G-50 pre-packed with 500 µl matrix	100 columns

Shipped at RT; store at 4 °C



## References

Judith M. Boer, Wolfgang K. Huber, Holger Sültmann, Friederike Wilmer, Anja von Heydebreck, Stefan Haas, Bernhard Korn, Bastian Gunawan, Andreas Vente, Laszlo Füzesi, Martin Vingron, and Annemarie Poustka Identification and Classification of Differentially Expressed Genes in Renal Cell Carcinoma by Expression Profiling on a Global Human 31,500-Element cDNA Array Genome Res., Nov 2001; 11: 1861 - 1870.

Ralf Ehricht, Thomas Kirner, Thomas Ellinger, Petra Foerster, and John S. McCaskill Monitoring the amplification of CATCH, a 3SR based cooperatively coupled isothermal amplification system, by fluorimetric methods Nucleic Acids Res., Nov 1997; 25: 4697 - 4699.

Federica Marini and Richard D. Wood A Human DNA Helicase Homologous to the DNA Cross-link Sensitivity Protein Mus308 J. Biol. Chem., Mar 2002; 277: 8716 - 8723.

Stefan Porubsky, Holger Schmid, Mahnaz Bonrouhi, Matthias Kretzler, Ernst Malle, Peter J. Nelson, and Hermann-Josef Gröne Influence of Native and Hypochlorite-Modified Low-Density Lipoprotein on Gene Expression in Human Proximal Tubular Epithelium Am. J. Pathol., Jun 2004; 164: 2175 - 2187.





# Chapter IV

## Compact Reaction Columns (CRC) Mobicols with Enzymatically Active Matrices

CRC with Immobilized Proteases

CRC with Immobilized Nucleases

CRC with other Immobilized Enzymes

Custom Immobilization



## Immobilized Enzymes for solid Phase Enzyme Reactions

Compact Reaction Columns (CRC) are small volume columns (Mobicols) containing a matrix with covalently immobilized enzymes. The patented immobilization chemistry was developed in cooperation with the Max Planck Society. It enables us to offer enzymes bound to F7m and G3m matrices with high enzymatic densities. The high enzyme densities of the CRCs result in fast substrate to product turnover. Enzyme reaction occurs when the substrate is loaded onto the column. The sample is recovered from the column quantitatively either by elution or by centrifugation. Since the enzymes are covalently bound, they remain in the column after reaction and product elution, making the CRC reusable. The eluted product solution is free of enzyme and does not require enzyme removal steps. This results in shorter lab protocols and handling procedures.

### Features

- Covalently bound enzymes
- No product contamination
- Suitable for small and large volumes
- Luer-lock connection to reservoir possible
- Short reaction protocol
- CRC is reusable
- Simple product recovery by centrifugation or elution
- Convenient washing enabled/possible
- High enzymatic densities

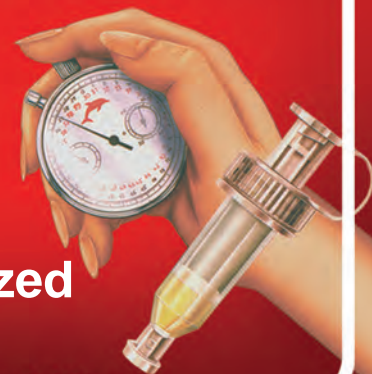
### Product Description

The immobilized enzymes are stable in aqueous media at a pH range of 5 to 10 and column bleeding is negligible. The „stiff“ linkers, which keep the enzymes from the matrix surface, effectively eliminate steric hindrance. This results in high activity of enzymes in the immobilized state. Column elution characteristics vary depending on the nature of the matrix used for immobilization.

**Cut down time  
and costs**

**with  
immobilized  
enzymes**

**MoBiTec**



- Columns with enzymes immobilized on matrix F7m are designed for applications involving either large substrate quantities, large volumes and/or large molecules.
- Columns with enzymes immobilized on G3m matrix are designed for applications involving small substrate quantities, small volumes and/or small molecules. Samples with a molecular weight  $\geq 10^3$  Daltons are recovered in the void volume when loaded on these columns.

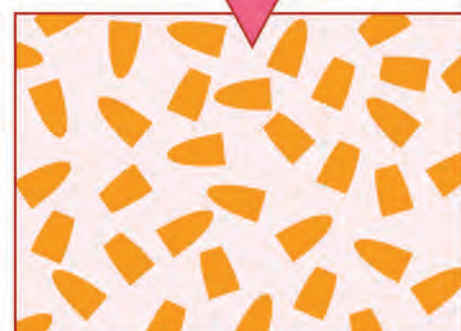
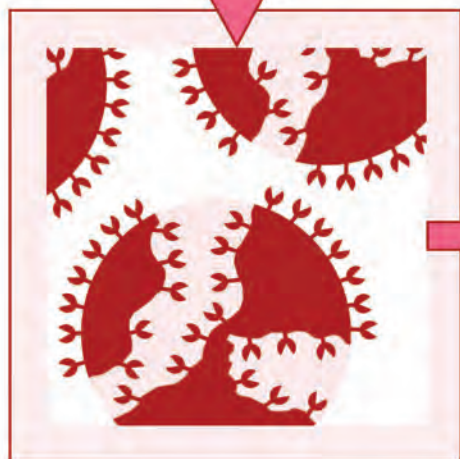
This well established technology allows you to expose your reaction solutions to very high concentrations of modifying enzymes. The exposure to high enzyme concentrations allows very short reaction times.

### Compact Reaction Columns (CRC)

Immobilized enzymes are supplied in versatile compact reaction columns (CRC) which fit into 1.5 or 2 ml microcentrifuge tubes. The columns have Luer-lock fittings, allowing direct syringe application of substrate solution, continuous flow processing of bulk solutions, or application of pressure for recovering the substrate. For small substrate volumes (approx. 50  $\mu$ l or less), most enzyme columns can be spun dry in benchtop centrifuges for fast, effective recovery.

**Easy handling, brief procedure in 4 steps**

- Equilibrate column with reaction buffer by flow-through (since column is provided with storage buffer)
- Load substrate solution in reaction buffer
- Incubate
- Elute product solution by:
  - a) centrifugation (smaller volumes, matrix G3m)
  - b) continuous flow mode (larger volumes, matrix F7m)

**Advantages**

- Very high enzyme activities
- No enzyme removal
- G3m: no sample dilution
- No long lasting gravity flow
- No chromatography system
- Easy handling, multiple sample processing in a few minutes



### Matrix Characteristics

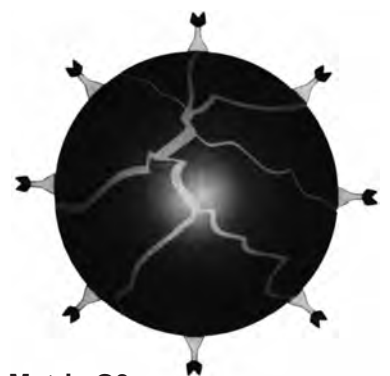
Two different immobilization matrices are available: F7m and G3m. They have different pore characteristics, as described below.



**F7m**  
Large pores-  
high activity!

### Matrix F7m

Matrix F7m has large pores. Molecules with up to  $10^7$  Dalton molecular weight (most enzymes and substrates) can enter these pores. The total surface of the material (including the surface inside the pores) is very large, resulting in an extremely high enzyme activity on the matrix. The product, which also enters the pores, has to be washed out of the matrix, leading to a dilution of the sample solution by additional 200  $\mu$ l to 500  $\mu$ l, depending on the substrate. If a dilution of the sample solution by 200  $\mu$ l or more cannot be tolerated, matrix G3m should be used.



**G3m**  
Small pores-  
no sample dilution!

### Matrix G3m

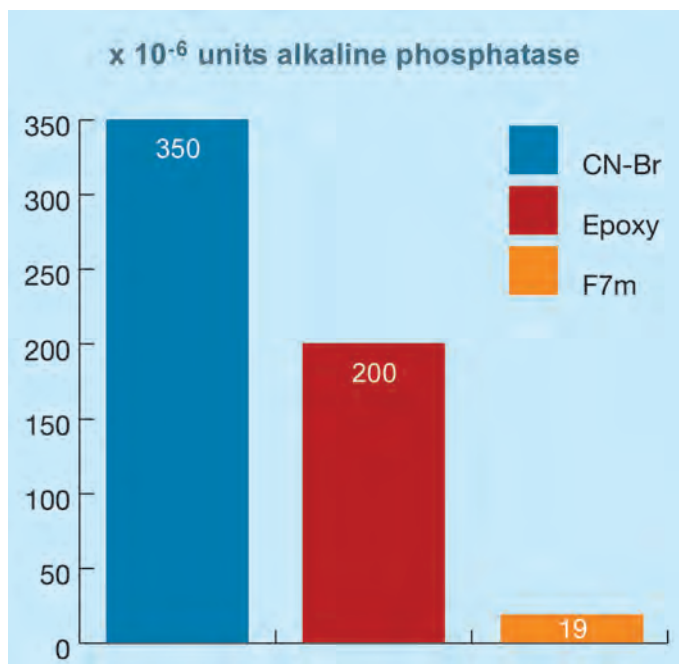
G3m has small pores; this results in excellent recovery characteristics (i.e. complete recovery in very small volumes). Molecules larger than  $10^3$  Dalton molecular weight (larger peptides, proteins and nucleic acids) cannot enter these pores. The total surface area of the material (only the outside of the spheres) is smaller than for F7m resulting in a smaller enzyme activity on the matrix. The immobilized enzymes as well as most substrates reside only in the volume between the matrix beads (in the dead volume). Thus, the sample solution can be completely removed by centrifugation without dilution. The design of the CR columns makes centrifugation handling easy. Sample solutions down to 20  $\mu$ l can be treated and recovered.

### Immobilization Chemistry

The immobilization chemistry has the particular advantage of being extremely stable in aqueous solution in the pH range 5 to 10. Bleeding-out is negligible. The rather stiff linkers separate the enzymes from the matrix surface thus eliminating steric hindrance. In the immobilized state the enzymes retain a high activity.

### Bleeding out in Comparison to Other Immobilization Chemistries

1,000 units alkaline phosphatase were immobilized on the same volume of different activated resins. 200  $\mu$ l Tris/HCl buffer pH 8.0 were applied on the columns for 20 minutes and recovered. The activity of the alkaline phosphatase in the sample was assayed. The relative amounts are shown in the figure below.



Comparison of three different matrices. According to these results, F7m is the matrix of choice, since bleeding out is significantly reduced compared to CNBr or Epoxy matrices.

### Columns, Filters and Buffers

Our 1 ml Mobicol column is specifically designed for this purpose. The enzymes are immobilized on 200  $\mu$ l matrix which is placed between two filters. The lower filters in the G3m columns have a pore size of 35  $\mu$ m, in F7m columns the lower filters have 10  $\mu$ m pore size. All upper filters have 90  $\mu$ m pore size. The filter material is of polyethylene. The enzyme matrix is delivered in storage buffer. Reaction, washing and storage buffers are provided for the first applications. All buffers are specified on the detailed data sheets provided with the columns.

The amount of liquid, which can be spun out of the CR-column to be replaced by new solution, is 90  $\mu$ l for F7m matrix and 76  $\mu$ l for G3m matrix.

### Effect of pH

The protein coupling to the matrix is stable in the range of pH 5 to pH 10. For short time periods (up to a few hours) the columns can be used also in the range pH 4 to pH 11. The immobilized enzymes should not, however, be stored below pH 5 or above pH 10. At extreme pH values the enzyme/matrix binding will degenerate.

### Biological Activity

DNA plasmids are biologically as active (in transformation) after passage through a CR-column (Proteinase K, RNase A) as the untreated molecules.



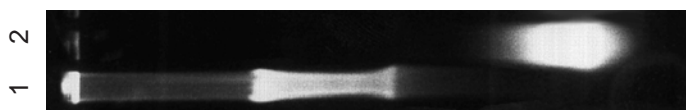
## Applications

### Removal of genomic DNA

Immobilized DNase I on G3m matrix is a useful tool for the removal of contaminating genomic DNA during RNA purification for single-cell cDNA-PCR: Standard DNase I digestion of RNA prepared from a single cell often leads to loss of material due to the required extraction step. Therefore, a method eliminating phenol extraction is extremely helpful. Using immobilized DNase I instead of soluble enzyme allows the efficient amplification of cDNA by PCR at the single-cell level in the absence of contaminating DNA.

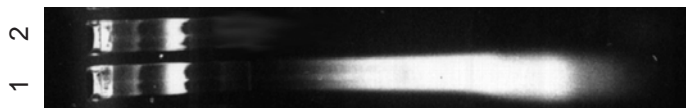
PCR is a process covered by patents owned by Hoffmann La-Roche.

### DNA digestion in a column



Immobilized DNase I on matrix G3m Agarose gel showing DNA before (lane 1) and after (lane 2) treatment on a DNase I column. The DNA is completely digested on the column.

### Easy digesting RNA during plasmid preps without phenol extractions

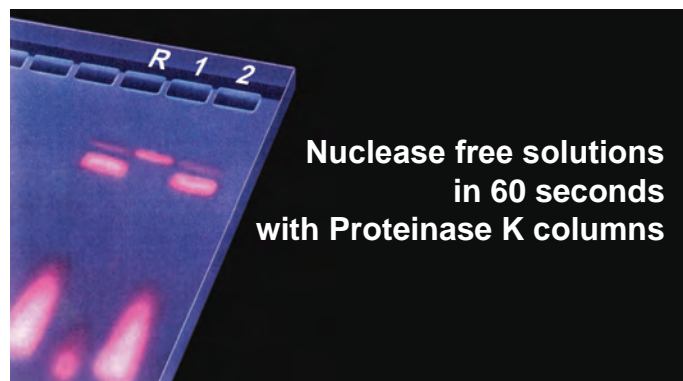


supercoiled plasmid DNA RNA

Agarose gel after separation of a clear lysate solution of a plasmid preparation containing DNA (supercoiled and nicked) and RNA (lane 1). Lane 2: Same solution after passage through a RNase G3m column. While the DNA remains unchanged the RNA is digested and cannot be detected.

### Linear vector DNA dephosphorylation using an alkaline phosphatase G3m CR-column

When cloning DNA fragments into plasmids such as expression vectors (i.e. plasmids up to 10 kb), the background resulting from recircularized plasmid vectors has to be minimized. This can be done by an alkaline phosphatase treatment of the linearized plasmid. Both ends will be dephosphorylated, thus preventing recircularization. The DNA fragments to be cloned, generated by restriction enzyme digestion, will provide the 5' terminal phosphate and thus ligate efficiently to the vector. Following plasmid dephosphorylation, the phosphatase activity must be removed before ligation in order to prevent dephosphorylation of the insert. To overcome this problem CR-columns with immobilized alkaline phosphatase provide dephosphorylated DNA ready for ligation without phosphatase activity contamination.

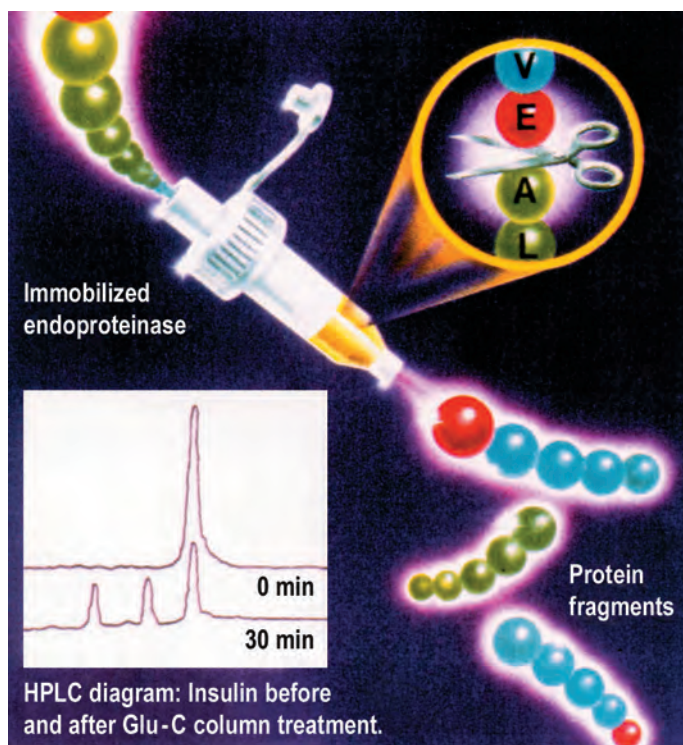


Nuclease free solutions in 60 seconds with Proteinase K columns

A DNA and RNA solution (reference, lane R) is contaminated with nucleases. After passage through a Proteinase K CR column, both solutions are incubated at 37°C for one hour, followed by agarose gel analysis. Untreated solution is digested (lane 1) while the recovered sample from the column (lane 2) is free of degradation.

### Insulin Box digestion by an endoproteinase Glu-C Compact Reaction Column (matrix G3m)

Since the endoproteinase is immobilized, it is completely retained in the column. Thus, the cleaved protein is entirely free of proteinases (and protease fragments); there is no need for further purification of the product.



HPLC diagram: Insulin before and after Glu-C column treatment.

Schematic drawing of proteins cleaved within a compact reaction column containing endoproteinase Glu-C immobilized on matrix G3m.

Small insert: 30 µg of insulin Box were loaded on an endoproteinase Glu-C compact reaction column with matrix G3m. The substrate solution was left in the column for 30 minutes and was analyzed by reverse phase HPLC. Peptides were detected at wavelength 215 nm.



### Order Information, Shipping & Storage

Order No.	Description	Amount
<b>Immobilized Proteinases (in CRC)</b>		
<b>P5101</b>	Endoproteinase Glu-C (Prot.V8) (one column with 200 µl matrix F7m)	900 U
<b>P3102</b>	Endoproteinase Glu-C (Prot.V8) (one column with 200µl matrix G3m)	22 U
<b>P5401</b>	Papain (one column with 200 µl matrix F7m)	23 U
<b>P3402</b>	Papain (one column with 200 µl matrix G3m)	0.6 U
<b>P5121</b>	Pepsin (one column with 200 µl matrix F7m)	16 mAnson U
<b>P3122</b>	Pepsin (one column with 200 µl matrix G3m)	0.4 mAnson U
<b>P5501</b>	Proteinase K (one column with 200 µl matrix F7m)	27 mAnson U
<b>P3502</b>	Proteinase K (one column with 200 µl matrix G3m)	0.7 mAnson U
<b>P5301</b>	TLCK-α -Chymotrypsin (one column with 200 µl matrix F7m)	55 U
<b>P3302</b>	TLCK-α -Chymotrypsin (one column with 200 µl matrix G3m)	1.4 U
<b>P5701</b>	TPCK-Trypsin (one column with 200 µl matrix F7m)	10,200 St-U
<b>P3702</b>	TPCK-Trypsin (one column with 200 µl matrix G3m)	260 St-U

### Immobilized Nucleases in CRC (except N3403)

<b>N5401</b>	DNase I (one column with 200 µl matrix F7m)	3,500 U
<b>N3402</b>	DNase I (one column with 200 µl matrix G3m)	88 U
<b>N3403</b>	DNase I (kit with 200 µl matrix and 5 empty columns)	88 U
<b>N5101</b>	RNase A (one column with 200 µl matrix F7m)	50 Kunitz-U
<b>N3102</b>	RNase A (one column with 200 µl matrix G3m)	2.5 Kunitz-U

### Immobilized Other Enzymes (in CRC)

<b>A5201</b>	Alkaline phosphatase (CIP) (one column with 200 µl matrix F7m)	1,000 U
<b>A3202</b>	Alkaline phosphatase (CIP) (one column with 200 µl matrix G3m)	100 U
<b>A5101</b>	β-Galactosidase (one column with 200 µl matrix F7m)	600 U
<b>A3102</b>	β-Galactosidase (one column with 200 µl matrix G3m)	15 U

shipped at RT; store at 4°C

**Note: Never freeze a CR-column!**

**A detailed handbook including all protocols is provided with the product. It is also available for download on our website [www.mobitec.com](http://www.mobitec.com).**

### All Columns are delivered with

- 200 µl immobilized enzyme matrix
- concentrated buffers for the first applications
- our CRC handbook about immobilized enzymes
- a data sheet with the description of the immobilized enzyme, the buffer composition and the protocol for the use of the column.



## References

William N. Addison, Fereshteh Azari, Esben S. Sørensen, Mari T. Kaartinen, and Marc D. McKee Pyrophosphate Inhibits Mineralization of Osteoblast Cultures by Binding to Mineral, Up-regulating Osteopontin, and Inhibiting Alkaline Phosphatase Activity *J. Biol. Chem.*, May 2007; 282: 15872 - 15883.

Jae Min Chong and David W. Speicher Determination of Disulfide Bond Assignments and N-Glycosylation Sites of the Human Gastrointestinal Carcinoma Antigen GA733-2 (CO17-1A, EGP, KS1-4, KSA, and Ep-CAM) *J. Biol. Chem.*, Feb 2001; 276: 5804 - 5813.

Scott Ficarro, Olga Chertihin, V. Anne Westbrook, Forest White, Friederike Jayes, Petr Kalab, Jarrod A. Marto, Jeffrey Shabanowitz, John C. Herr, Donald F. Hunt, and Pablo E. Visconti Phosphoproteome Analysis of Capacitated Human Sperm. EVIDENCE OF TYROSINE PHOSPHORYLATION OF A KINASE-ANCHORING PROTEIN 3 AND VALOSIN-CONTAINING PROTEIN/p97 DURING CAPACITATION *J. Biol. Chem.*, Mar 2003; 278: 11579 - 11589.

Marcin Gierdalski and Sharon L. Juliano Factors Affecting the Morphology of Radial Glia, Cereb Cortex, Jun 2003; 13: 572 - 579.

Jeff Holderness, Larissa Jackiw, Emily Kimmel, Hannah Kerns, Miranda Radke, Jodi F. Hedges, Charles Petrie, Patrick McCurley, Pati M. Glee, Aiyappa Palecanda, and Mark A. Jutila Select Plant Tannins Induce IL-2R $\alpha$  Up-Regulation and Augment Cell Division in  $\gamma\delta$  T Cells *J. Immunol.*, Nov 2007; 179: 6468 - 6478.

R.E. Leube, U. Leimer, C. Grund, W.W. Franke, N. Harth, and B. Wiedenmann Sorting of synaptophysin into special vesicles in nonneuroendocrine epithelial cells *J. Cell Biol.*, Dec 1994; 127: 1589.

Thomas S. Nühse, Thomas Boller, and Scott C. Peck A Plasma Membrane Syntaxin Is Phosphorylated in Response to the Bacterial Elicitor Flagellin *J. Biol. Chem.*, Nov 2003; 278: 45248 - 45254.

Thomas S. Nühse, Allan Stensballe, Ole N. Jensen, and Scott C. Peck Large-scale Analysis of in Vivo Phosphorylated Membrane Proteins by Immobilized Metal Ion Affinity Chromatography and Mass Spectrometry *Mol. Cell. Proteomics*, Nov 2003; 2: 1234.

A. Pries, H. Priefert, N. Krüger, and A Steinbüchel.. Identification and characterization of two *Alcaligenes eutrophus* gene loci relevant to the poly( $\beta$ -hydroxybutyric acid)-leaky phenotype which exhibit homology to *ptsH* and *ptsI* of *Escherichia coli*. *J. Bacteriol.*, Sep 1991; 173: 5843 - 5853.

J. Ronnenberg et al. Protein Sequenzierung mit immobilisierten sequenz-spezifischen Endoproteinasen

J. Ronnenberg, B. Preitz, G. Wostemeier, and S. Diekmann Immobilized residue-specific endoproteinasen for protein sequencing. *J Chromatogr B Biomed Appl*, Jun 1994; 656(1): 169-77.

N. M. Sawtell, D. K. Poon, C. S. Tansky, and R. L. Thompson The Latent Herpes Simplex Virus Type 1 Genome Copy Number in Individual Neurons Is Virus Strain Specific and Correlates with Reactivation *J. Virol.*, Jul 1998; 72: 5343 - 5350.

Richard L. Thompson and N. M. Sawtell PATHOGENESIS AND IMMUNITY: Replication of Herpes Simplex Virus Type 1 within Trigeminal Ganglia Is Required for High Frequency but Not High Viral Genome Copy Number Latency *J. Virol.*, Jan 2000; 74: 965 - 974.

R. L. Thompson and N. M. Sawtell PATHOGENESIS AND IMMUNITY: Herpes Simplex Virus Type 1 Latency-Associated Transcript Gene Promotes Neuronal Survival *J. Virol.*, Jul 2001; 75: 6660 - 6675.





# Mobicols

SEPARATION

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ISOLATION

ISOLATION

TREATMENT

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PURIFICATION

PURIFICATION

**ITALY :**

**Duotech** srl

**Ph :** 02.33106630

**fax :** 02.33106640

**e-mail :** info@duotech.ir

**internet :** www.duotech.it

**[www.mobitec.com](http://www.mobitec.com)**

MoBiTec GmbH,

Lotzestrasse 22a, 37083 Goettingen, Germany

Phone: +49 (0)551-70722-0

Fax: +49 (0)551-70722-22

E-mail: info@mobitec.com



**Mo Bi Tec**  
MOLECULAR BIOTECHNOLOGY