OFNEDAL TEOLINOLOG			
GENERAL TECHNOLOG	Product/activity	Location	URL
Company	_	Location	ONL
Advion BioSciences	Has an ESI chip for MS manufactured by Intellisense.		
	Contract LC company.	Ithaca, New York	www.advion.com
Affibody	Uses propietary protein to study protein-protein		
4 D.D. 4	interactions. Associated with Gyros.	Stockholm, Sweden	www.affibody.com
ARRM	Robotics, associated with Hochstrasser and Genetix.	Adelaide, Australia	www.arrm.com
Biacore	Ligand fishing; surface plasmon resonance		
	for real-time detection and monitoring of		
B: B ! !!	biomolecular binding events.	Uppsala, Sweden	www.biacore.se/proteomics
BioRobotics	Robotics for arrays and colony picking.	Cambridge, UK	www.biorobotics.com
Borealis Biosciences	Protein identification, affinity chromatography.	Toronto, Canada	www.protein-affinity.com
Caliper	Microfluidics-based 'lab-on-a-chip'. Capillary-based		
	'sipper chip' for continuous screening assays.		
	Fluorogenic, electrophoretic mobility, including		
	kinases and phosphatases. To come are calcium		
	flux and membrane potential. Collaborates with	Manufactura Villago California	
Convious Dhoumassou tisals	Structural GenomiX and Agilent.	Mountain View, California	www.calipertech.com
Caprion Pharmaceuticals	CellCarta organelle purification-based identification	Mantard Const.	
Cinhagan	of rare proteins.	Montreal, Canada	www.caprion.com
Ciphergen	Protein-chip reader, chip software, chip arrays		
	(hydrophilic/hydrophobic, anion/cation exchange,		
	immobilized metal), SELDI chip, protein-chip	Exament California	vereze sinhouses som
Cutomic	arrays for affinity capture.	Fremont, California	www.ciphergen.com
Cytomyx	Provides proteomics services from sample	Cambridge LIV	NAMES OF THE PARTY COME
Europroteomo	preparation through to MS. Works on epithelial cancers.	Cambridge, UK Hennigsdorf, Germany	www.cytomyx.com
Europroteome	GelPix spot excision, array products, robotics.		www.europroteome.com
Genetix		New Milton, UK	www.genetix.com
Genomic Solutions	Investigator proteomic system, fully integrated solution for identifying and characterizing proteins.		
	ProGest sample preparation station.	Ann Anhan Mishiann	
Chron	1 1 1	Ann Arbor, Michigan	www.genomicsolutions.com
Gyros	Nanoscale 'lab-on-a-CD' for MS.	Uppsala, Sweden	www.gyrosmicro.com
Ingeny	On the verge of introducing an automated	Coos Noth orlando	
Integrative Protection	2D-gel system.	Goes, Netherlands	www.ingeny.com
Integrative Proteomics	Uses Bruker XR, NMR, MS equipment for	Towarta Canada	
Lorgo Coolo Diology	functional proteomics.	Toronto, Canada	www.integrativeproteomics.com
Large Scale Biology	Produces therapeutic proteins in plants. Is		
	developing an antibody chip with Biosite	Vacavilla California	www.lsbc.com
LumiOuto	2D-gel factory.	Vacaville, California	
LumiCyte MediChem	Protein biomarker profiles for drug development.  Protein expression, crystallization, structure,	Fremont, California	www.lumicyte.com
Medicilem	through subsidiary companies.	Woodridge Illinois	www.medichem.com/
	unrough subsidiary companies.	Woodridge, Illinois	services/proteomics.htm
Myriad Genetics	ProNet yeast two-hybrid protein interaction		services/proteonnes.nun
Myriad defletics	system, ProSpec MS-based drug discovery.	Salt Lake City, Utah	www.myriad.com
NextGen Sciences	Platform technologies in proteomics, transcriptomics	Sait Lake City, Otali	www.myriad.com
NextGerr Sciences	and genomics.	Huntingdon, UK	www.nextgensciences.com
Oxford GlycoSciences	2D gels, developing ICAT/MALDI-TOF system.	Oxford, UK	www.ogs.com
Pepscan Systems	Protein–protein interactions based on epitopes.	Oxioiu, OK	www.ogs.com
r cpscarr bysterns	epitope mapping, combinatorial peptide arrays and		
	peptidomimetics. Lead development service.	Lelystad, Netherlands	www.pepscan.nl
Phylos	Antibody-based high-throughput 'HIP' chip;	Letystad, Netherlands	www.pepscan.m
Tiylos	ProFusion selection technology based on protein		
	fused to its own mRNA.	Lexington, Massachusetts	www.phylos.com
Protagen	Phosphorylation analysis.	Bochum, Germany	www.prytos.com www.protagen.de
Proteomic Solutions	Proteomics service company.	St Marcel, France	www.protagen.de www.proteomicsolutions.fr
Protein Pathways	Pathway analysis in mammals and microbes.	Los Angeles	www.proteinpathways.com
Proteome Sciences	Identifies protein markers in human disease	Los migeres	
	targets. US subsidiary is Intronn LLC.	Cobham, Surrey, UK	www.proteome.co.uk
Proteome Systems	Glycosylation analysis, 2D-gel excision equipment,	Coonam, Suriey, UK	" " W.P. OtCOINE.CO.UK
. Totoomo Gyatema	chemical printer for electroblotting gels to		
	membranes, Axima CFR MS machine.	Sydney, Australia	www.nroteomesystems.com
ProteoSys		Sydney, Australia	www.proteomesystems.com
Toteosys	Analyses modifications, quantifies down to	Mainz Carmany	unun protoccue com
Rigel	sub-attomolar range. Oncology proteins, ubiquitin ligase.	Mainz, Germany South San Francisco, CA	www.proteosys.com www.rigel.com
Roche	Proteomics research centre.	Basel, Switzerland	www.rigei.com www.roche.com
Sense Proteomic	COVET functional protein array.	Cambridge, UK	www.sensetherapeutic.com
Sensium Technologies	Uses biosensors to detect low abundance proteins,	Cambridge, OK	www.sensemerapeunc.com
Solisian reciniologies	proteomics subsidiary of Aurora/Helix BioPharma.	Edmonton, Canada	www.heliybionharma.com
Somal ogic		Boulder, Colorado	www.helixbiopharma.com
SomaLogic Syrry	Aptamer arrays, partner with Celera. Structural proteomics, proteomics discovery	Douidel, Colorado	www.somalogic.com
Syrrx		San Diego California	MANAGEMENT COM
Virtek	platform with Eli Lilly.  Deskton array reader can be used for 2D gels	San Diego, California Ontario, Canada	www.syrrx.com www.virtek.ca
WITA Proteomics	Desktop array reader can be used for 2D gels.	Omario, Canada	www.vii tek.ca
VVITATIOLEOHIICS	High-resolution 2D-gel platform, protein identification.	Tetlow Carmany	unun wita_protoomics_com
Zymark	Alliance with Pharmagene.  Robots for screening, liquid handling, plate management.	Tetlow, Germany Hopkinton, Massachusetts	www.wita-proteomics.com
Zymark Zyomyx		Hayward, California	www.zymark.com
ZyUITIYX	Protein-chip development.	riaywaru, Calliofilia	www.zyomyx.com

MASS SPECTROMETRY			
Company	Products	Location	URL
Agilent Technologies	Ion-trap MS machines.	Palo Alto, California	www.agilent.com/chem
Amersham Biosciences	Ettan DALT II MALDI-TOF MS.	Uppsala, Sweden	www.amershambiosciences.com
Analytica of Branford	Prototype MS developer.	Branford, Connecticut	www.aob.com
Applied Biosystems	API series (with MDS Sciex) and Voyager		
	MS machines.	Foster City, California	www.appliedbiosystems.com
Bio-Rad	ProteomeWorks System, with MicroMass.	Hercules, California	www.
210 1144	Trocome works system, with merowass.	Trereures, Gamorina	proteomeworkssystem.com
Bruker Daltonics	Lilendary 2000 TOP/TOP to a low MC CNAD		proteomeworkssystem.com
Bruker Daitoriics	Ultraflex 3000 TOF/TOF tandem MS, SNAP,	D C	11 1 1
1.19	biotools, ProteinScape software.	Bremen, Germany	www.daltonics.bruker.com
Hitachi Instruments	Many kinds of MS, LC & analytical equipment.	Yokohama, Japan	www.hii-hitachi.com
MDS Sciex	API series LC/MS systems.	Toronto, Canada	www.sciex.com
Micromass	ProteomeWorks System marketed by Bio-Rad.	Manchester, UK	www.micromass.co.uk
ThermoFinnigan	Quadrupole, TOF and ion-trap MS machines.	San Jose, California	www.thermofinnigan.com
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REAGENT SUPPLIERS			
Company	Products	Location	URL
Amicon	Immobilon transfer membranes division of Millipore.	Location	www.millipore.com/lifescience
ATTICOT	immobilion transfer memoranes division of willipore.	D 10 1 M 1 "	
		Bedford, Massachusetts	/products.nsf/docs/amicon
BioWhittaker	Reagents, part of Cambrex Life Sciences. Has lab		
	service centre in North Brunswick, New Jersey.	East Rutherford, New Jersey	www.cambrex.com
CBS Scientific	Reagents for gels.	Del Mar, California	www.cbssci.com
Crescent Chemical	Electrophoresis and LC chemicals.	New York	www.creschem.com
Fermentas	Reagents, protein standards.	Vilnius, Lithuania	www.fermentas.com
IBA	Protein expression and purification with STREP-tag	Desired	
	technology.	Cättingen Commeny	www.iba-go.de
IONI Diama dia ala		Göttingen, Germany	
ICN Biomedicals	Reagents, radioisotopes.	Costa Mesa, California	www.icnbiomed.com
Invitrogen	Reagents, membranes, filters.	Paisley, UK	www.invitrogen.com
Jule Biotechnologies	Precast gels.		hometown.aol.com/
		New Haven, Connecticut	precastgel/index.htm
Molecular Probes	Stains.	Leiden, Netherlands	www.probes.com
Novagen	Protein purification and analysis products.	Madison, Wisconsin	www.novagen.com
Pierce	Chemicals, reagents, trypsin, antibodies, cytokines.	Rockford, Illinois	www.piercenet.com
	Expression cloning system.		•
Promega	Expression cioning system.	Madison, Wisconsin	www.promega.com
Qiagen	Protein expression and purification, also has		
	Biorobot 8000 for purifying proteins.	Hilden, Germany	www.qiagen.com
R. Shadel	Gel boxes, combs, gaskets, spacers.	San Francisco	www.shadel.com
Schleicher & Schuell	Filters, membranes.	Dassel, Germany	www.s-und-s.de
Sigma-Aldrich	Reagents, purified trypsin.	St Louis, Missouri	www.sigma-aldrich.com
Stratagene	In vitro mutagenesis kit.	La Jolla, California	www.stratagene.com
Zaxis	Precast gels.	Hudson, Ohio	www.zaxis-inc.com
Laxio	recust gets.	Tradon, omo	WWW.Zukio Incicom
SOFTWARE			
	Duaduata	Leastion	LIDI
Company	Products	Location	URL
Accelrys	Gene Atlas, Atlas Base.	San Diego, California	www.accelrys.com
	GLD ligand identification, ProChart		
	pathway identification.	Newtown, Pennsylvania	www.axcellbio.com
AxCell Biosciences	pathway identification.	Newtown, Pennsylvania	
AxCell Biosciences BIND	pathway identification. Database of pathways and interactions.	Newtown, Pennsylvania	www.axcellbio.com Toronto, Canada www.bind.ca
AxCell Biosciences BIND	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec),	Newtown, Pennsylvania	
AxCell Biosciences BIND	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date	,	Toronto, Canada www.bind.ca
AxCell Biosciences BIND BioBridge Computing	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.	Newtown, Pennsylvania Lund, Sweden	
AxCell Biosciences BIND BioBridge Computing	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent	,	Toronto, Canada www.bind.ca
AxCell Biosciences BIND BioBridge Computing	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural	,	Toronto, Canada www.bind.ca
AxCell Biosciences BIND BioBridge Computing	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by	,	Toronto, Canada www.bind.ca
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AxCell Biosciences BIND BioBridge Computing Cognia	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark.  Z3 @d-gel image analysis system, Protocall MS	Lund, Sweden  New York	Toronto, Canada www.bind.ca www.biobridge.se www.cognia.com
AxCell Biosciences  BIND BioBridge Computing  Cognia  Compugen	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark.  Z3 @d-gel image analysis system, Protocall MS analysis tools.	Lund, Sweden	Toronto, Canada www.bind.ca www.biobridge.se
AxCell Biosciences  BIND BioBridge Computing  Cognia  Compugen	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark.  Z3 @d-gel image analysis system, Protocall MS analysis tools.  DeltaBase, functional annotation DB,	Lund, Sweden  New York  Tel Aviv, Israel	Toronto, Canada www.bind.ca www.biobridge.se www.cognia.com www.cgen.com
AxCell Biosciences  BIND BioBridge Computing  Cognia  Compugen  Deltagen	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark. Z3 @d-gel image analysis system, Protocall MS analysis tools.  DeltaBase, functional annotation DB, phentoype and knockout information.	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California	Toronto, Canada www.bind.ca www.biobridge.se www.cognia.com www.cgen.com
AxCell Biosciences  BIND  BioBridge Computing  Cognia  Compugen  Deltagen  Hybrigenics	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark. Z3 @d-gel image analysis system, Protocall MS analysis tools.  DeltaBase, functional annotation DB, phentoype and knockout information. PIMRider, protein pathways and interactions software.	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California Paris, France	Toronto, Canada www.bind.ca www.biobridge.se www.cognia.com www.cgen.com www.deltagen.com pim.hybrigenics.com
AxCell Biosciences  BIND  BioBridge Computing  Cognia  Compugen  Deltagen  Hybrigenics Incyte Genomics	pathway identification. Database of pathways and interactions. PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001. Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark. Z3 @d-gel image analysis system, Protocall MS analysis tools. DeltaBase, functional annotation DB, phentoype and knockout information. PIMRider, protein pathways and interactions software. Life Express protein database.	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California	Toronto, Canada www.bind.ca www.biobridge.se www.cognia.com www.cgen.com
AxCell Biosciences  BIND  BioBridge Computing  Cognia  Compugen  Deltagen  Hybrigenics Incyte Genomics	pathway identification. Database of pathways and interactions. PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001. Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark. Z3 @d-gel image analysis system, Protocall MS analysis tools. DeltaBase, functional annotation DB, phentoype and knockout information. PIMRider, protein pathways and interactions software. Life Express protein database. Human protein index, molecular anatomy	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California Paris, France	Toronto, Canada www.bind.ca www.biobridge.se www.cognia.com www.cgen.com www.deltagen.com pim.hybrigenics.com www.incyte.com
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AxCell Biosciences  BIND  BioBridge Computing  Cognia  Compugen  Deltagen  Hybrigenics Incyte Genomics Large Scale Biology	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark.  Z3 @d-gel image analysis system, Protocall MS analysis tools.  DeltaBase, functional annotation DB, phentoype and knockout information.  PIMRider, protein pathways and interactions software.  Life Express protein database.  Human protein index, molecular anatomy pathways, molecular effects of disease databases.	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California Paris, France Palo Alto, California	Toronto, Canada www.bind.ca www.biobridge.se www.cognia.com www.cgen.com www.deltagen.com pim.hybrigenics.com www.incyte.com
AxCell Biosciences  BIND  BioBridge Computing  Cognia  Compugen  Deltagen  Hybrigenics Incyte Genomics Large Scale Biology	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark.  Z3 @d-gel image analysis system, Protocall MS analysis tools.  DeltaBase, functional annotation DB, phentoype and knockout information.  PIMRider, protein pathways and interactions software. Life Express protein database.  Human protein index, molecular anatomy pathways, molecular effects of disease databases.  Mascot software searches sequence databases with	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California Paris, France Palo Alto, California  Germantown, Maryland	Toronto, Canada www.bind.ca www.biobridge.se  www.cognia.com www.cgen.com www.deltagen.com pim.hybrigenics.com www.incyte.com www.lsbc.com
AxCell Biosciences BIND BioBridge Computing  Cognia  Compugen Deltagen Hybrigenics Incyte Genomics Large Scale Biology  Matrix Science	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark.  Z3 @d-gel image analysis system, Protocall MS analysis tools.  DeltaBase, functional annotation DB, phentoype and knockout information.  PIMRider, protein pathways and interactions software. Life Express protein database.  Human protein index, molecular anatomy pathways, molecular effects of disease databases.  Mascot software searches sequence databases with MS signatures.	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California Paris, France Palo Alto, California  Germantown, Maryland  London, UK	Toronto, Canada www.bind.ca www.biobridge.se  www.cognia.com www.cognia.com www.deltagen.com pim.hybrigenics.com www.incyte.com www.lsbc.com
AxCell Biosciences BIND BioBridge Computing  Cognia  Compugen Deltagen Hybrigenics Incyte Genomics Large Scale Biology Matrix Science Media Cybernetics	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark.  Z3 @d-gel image analysis system, Protocall MS analysis tools.  DeltaBase, functional annotation DB, phentoype and knockout information.  PIMRider, protein pathways and interactions software.  Life Express protein database.  Human protein index, molecular anatomy pathways, molecular effects of disease databases.  Mascot software searches sequence databases with MS signatures.  ArrayPro image-analysis software.	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California Paris, France Palo Alto, California  Germantown, Maryland	Toronto, Canada www.bind.ca www.biobridge.se  www.cognia.com www.cgen.com www.deltagen.com pim.hybrigenics.com www.incyte.com www.lsbc.com
AxCell Biosciences  BIND BioBridge Computing  Cognia  Compugen  Deltagen  Hybrigenics Incyte Genomics Large Scale Biology  Matrix Science  Media Cybernetics	pathway identification.  Database of pathways and interactions.  PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001.  Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark.  Z3 @d-gel image analysis system, Protocall MS analysis tools.  DeltaBase, functional annotation DB, phentoype and knockout information.  PIMRider, protein pathways and interactions software.  Life Express protein database.  Human protein index, molecular anatomy pathways, molecular effects of disease databases.  Mascot software searches sequence databases with MS signatures.  ArrayPro image-analysis software.  ProteomeWorks, MassLynx, MetaboLynx	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California Paris, France Palo Alto, California  Germantown, Maryland  London, UK Silver Spring, Maryland	Toronto, Canada www.bind.ca www.biobridge.se  www.cognia.com www.cognia.com www.deltagen.com pim.hybrigenics.com www.incyte.com www.matrixscience.com www.matrixscience.com
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AxCell Biosciences BIND BioBridge Computing Cognia  Compugen Deltagen Hybrigenics Incyte Genomics Large Scale Biology Matrix Science Media Cybernetics Micromass Nonlinear Dynamics	pathway identification. Database of pathways and interactions. PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001. Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark. Z3 @d-gel image analysis system, Protocall MS analysis tools. DeltaBase, functional annotation DB, phentoype and knockout information. PIMRider, protein pathways and interactions software. Life Express protein database. Human protein index, molecular anatomy pathways, molecular effects of disease databases. Mascot software searches sequence databases with MS signatures. ArrayPro image-analysis software. ProteomeWorks, MassLynx, MetaboLynx MS software. Phoretix, Progenesis gel image-analysis software.	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California Paris, France Palo Alto, California  Germantown, Maryland  London, UK Silver Spring, Maryland  Manchester, UK Newcastle upon Tyne, UK	Toronto, Canada www.bind.ca www.biobridge.se  www.cognia.com www.cgen.com www.deltagen.com pim.hybrigenics.com www.incyte.com www.matrixscience.com www.mediacy.com www.micromass.co.uk www.nonlinear.com
AxCell Biosciences BIND BioBridge Computing Cognia  Compugen Deltagen Hybrigenics Incyte Genomics Large Scale Biology Matrix Science Media Cybernetics Micromass Nonlinear Dynamics	pathway identification. Database of pathways and interactions. PIUMS (Protein Identification Using Mass Spec), Pepex peak-picking software, launch date November 2001. Protein catabolism database (ubiquitin-dependent turnover for regulation of protein levels, structural homology elements for substrate recognition by E3 enzymes). Also markets Biobase, from University of Aarhus, Denmark. Z3 @d-gel image analysis system, Protocall MS analysis tools. DeltaBase, functional annotation DB, phentoype and knockout information. PIMRider, protein pathways and interactions software. Life Express protein database. Human protein index, molecular anatomy pathways, molecular effects of disease databases. Mascot software searches sequence databases with MS signatures. ArrayPro image-analysis software. ProteomeWorks, MassLynx, MetaboLynx MS software.	Lund, Sweden  New York  Tel Aviv, Israel  Redwood City, California Paris, France Palo Alto, California  Germantown, Maryland  London, UK Silver Spring, Maryland  Manchester, UK	Toronto, Canada www.bind.ca www.biobridge.se  www.cognia.com www.cgen.com www.deltagen.com pim.hybrigenics.com www.incyte.com www.matrixscience.com www.mediacy.com www.micromass.co.uk
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