

## Overview of the LiMB database

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John R. Lawton, Frances A. Martinez and Christian Burks\*

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Theoretical Biology and Biophysics Group, T-10, MS K710, Los Alamos National Laboratory, Los Alamos, NM 87545, USA

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### ABSTRACT

The rapidly increasing number of databases relevant to molecular biology has given rise to a need for a coordinated effort to identify, characterize, and link them. The LiMB database, which contains information about molecular biology and related databases, is a step in that direction. It serves molecular biologists seeking data sets containing information relevant to their research, and is also intended to anticipate the needs of database designers and managers building software links for related data sets. We present an abbreviated version of the database here; the full database is available free of charge as described below.

### INTRODUCTION

The rapidly increasing number of computer databases in the field of molecular biology has created a need for a systematic, coordinated approach to designing, developing, and maintaining these databases [1]. A prerequisite for much of this work is an overview of existing databases; until now, there has been no 'database of databases' that would support developing and maintaining such an overview. The LiMB (**L**isting of **M**olecular **B**iology **D**atabases) database was created to fill that gap.

The LiMB database is intended to facilitate locating and accessing databases relevant to molecular biology. It was begun in October 1986; the first (Release 1.0) formal release, containing information on 53 databases, was distributed in February 1988 [2]. It has been distributed directly to over 300 people, with a greater number of researchers accessing it through six network servers that have mounted copies of LiMB for the communities they serve.

### GENERAL DESCRIPTION

Each entry in LiMB currently consists of 54 fields, including: database staff names and addresses; database maintenance hardware and software; scope of coverage and database goals; details about submission and access to the data sets; database size; and types of data covered by the database. These data items are described in greater detail below. LiMB entries are based on questionnaires filled out by the database managers or, in the absence of a completed questionnaire, on secondary sources (e.g., a journal article). LiMB is currently maintained on a Sun 3/280 with UNIX system tools; in the near future it will be maintained using a relational DBMS.

LiMB serves both the potential users of these databases (by helping them find and contact the databases which contain the information they seek) and the designers and managers of the databases (by providing a centralized source of information that can aid coordination

## Nucleic Acids Research

**Table I.** LiMB data fields and definitions.

Field	Definition
entry	Name of entry
number	Accession number (will always be associated with a given data bank)
history	History of entry
status	Correspondence status
res.nam	Name of respondent
res.add	Address of respondent
res.tel	Telephone number of respondent (as dialed in the USA)
res.net	Network address of respondent
gen.nam	Name for general inquiry
gen.add	Address for general inquiry
gen.tel	Telephone number for general inquiry (as dialed in the USA)
gen.net	Network address for general inquiry
con.nam	Name for contributing data
con.add	Address for contributing data
con.tel	Telephone number for contributing data (as dialed in the USA)
con.net	Network address for contributing data
acc.nam	Name for acquiring data
acc.add	Address for acquiring data
acc.tel	Telephone number for acquiring data (as dialed in USA)
acc.net	Network address for acquiring data
name.now	Official name of database
name.alt	Alternative official/unofficial names of database
name.bad	Obsolete or incorrect names of database
source	Source of data in database
collab.in	Formal collaborators for collection of data
collab.out	Formal collaborators for distribution of data
funding	Funding base for data bank
citation	Recent publications describing database
charter	Formal or informal charter of data bank effort
cross	Other databases to which this database is explicitly cross-referenced
data.pri	Primary data items in the database
data.sec	Secondary data items in the database
hardware	Hardware used for maintaining the data
operate	Operating system for database hardware
dbms	Software system used for maintaining data
language	Programming language used for software system
format	Format used for distributed flat text files
software	Software distributed with the database
access	Limitations on access to data
update	Frequency of database updates
con.onl	Can contributions be made to database on-line?
con.mag	Can contributions be made to database on magnetic tape?
con.flp	Can contributions be made to database on floppy disk?
con.elm	Can contributions be made to database by electronic mail?
con.pap	Can contributions be made to database on hardcopy?
acc.onl	Is database distributed on-line?
acc.mag	Is database distributed on magnetic tape?
acc.flp	Is database distributed on floppy disk?
acc.elm	Is database distributed by electronic mail?
acc.pap	Is database distributed on hardcopy?
byt.all	Number of bytes contained in the database
byt.pri	Number of bytes contained in primary data items in the database
ent.pri	Number of entries in the database
comment	Information not contained in other fields
///	end of entry

between the various databases). It was begun as a simple directory; in the future it will expand its role as a centralized information resource by providing a platform for designing front-ends for automatic access to distributed biological data sets.

LiMB is available free of charge through one of the following media: electronic network mail, MS-DOS based floppy disk, or printed form. Requests for the database should include name, address (e-mail or postal service), and desired medium for transmission; these requests may be directed through postal service<sup>1</sup> or electronic network mail.<sup>2</sup>

The project is still in its formative stage. We would greatly appreciate any ideas for improving future releases and would also appreciate being informed of any molecular biology or related databases not listed in Release 1.0 that could be included in future releases.

## DATA FIELDS

Table I provides a listing of all the fields found—when the information is available and pertinent—in a flat file LiMB entry.

## THE DATA

A listing of all the LiMB entries (in abbreviated form) is presented in Appendix 1.0; an index based on the data items contained in the various databases is presented in Appendix 2.0. This represents an augmented and updated listing (Release 1.1) relative to Release 1.0 of LiMB [2] and earlier, preliminary versions of these data [3,4].

## ACKNOWLEDGEMENTS

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\*To whom correspondence should be addressed

<sup>1</sup> LiMB Database; T-10, MS K710; Los Alamos National Laboratory; Los Alamos, NM 87545; USA.

<sup>2</sup> limb@lanl.gov

## REFERENCES

1. Burks, C., Lawton, J.R. and Bell, G.I. (1988) *Science* **241**, 888.
2. Burks, C. and Lawton, J.R. (1988) LiMB database: Release 1.0. Unpublished release notes, Los Alamos National Laboratory, Los Alamos, NM, U.S.A..
3. Burks, C. (1987) In *Final Report for the EMBL/NIH Workshop on Future Databases for Molecular Biology*, European Molecular Biology Laboratory, Heidelberg (unpublished report).
4. von Heijne, G. (1987) 'Sequence Analysis in Molecular Biology,' Academic Press, New York, pp. 153–60.

**APPENDIX 1.0: ABBREVIATED LiMB ENTRIES**

entry	AAAAAAA (Name of entry.)	data.pri	[4.5S rRNA] nucleotide sequences; [5S rRNA] nucleotide sequences; [5.8S rRNA] nucleotide sequences.
name.now	Name of database.	data.sec	[nucleotide sequence, rRNA] functional features.
number	Number that will always be associated with a given data bank.	gen.nam	Dr. Volker Erdmann
data.pri	Primary data items in the database.	gen.add	Institut fur Biochemie
data.sec	Secondary data items in the database.		FB Chemie
gen.nam	Name for general inquiry.		Freie Universitat Berlin
gen.add	Address for general inquiry.		Otto-Hahn-Bau
gen.tel	Telephone number for general inquiry as dialed in U.S.A.)		Thielallee 63
gen.net	Network address for general inquiry.		D-1000 Berlin 33 (Dahlem)
///			WEST GERMANY
entry	AANSPII	gen.tel	[011-49] 30-838 6002
name.now	Amino Acid and Nucleotide Sequences of Proteins of Immunological Interest	///	
number	10001	entry	CARBBANK
data.pri	[immunoglobulin] amino acid sequences; [immunoglobulin] nucleotide sequences; sequence alignments.	name.now	CarbBank
data.sec	[amino acid sequence, immunoglobulin] functional features; [nucleotide sequence, immunoglobulin] functional features.	number	10003
gen.nam	Dr. Elvin Kabat	data.pri	carbohydrate sequences.
gen.add	National Institutes of Health Bldg. 8 Room 126 Bethesda, MD 20892 U.S.A.	data.sec	[carbohydrate sequence] functional features; index terms; literature citations; [carbohydrate sequence] sources.
gen.tel	301-496-0316	gen.nam	Dr. Scott Doubet
///		gen.add	Complex Carbohydrate Research Center
entry	AMINODB		Russell Lab
name.now	Amino Acid Database		P.O. Box 5677
number	10042		Athens, GA 30613
data.pri	[amino acid] molecular properties.		U.S.A.
data.sec	[molecular biology] literature citations.	gen.tel	404-546-3555
gen.nam	Dr. Cornelius Froemmel	gen.net	carbbank@uga.bitnet
gen.add	DDR-1040 Berlin Institut fur Biochemie Hessische Strasse 3-4 EAST GERMANY	///	
///		entry	CAS
entry	BIOSISP	name.now	CAS ONLINE
name.now	BIOSIS Previews	number	10030
number	10031	data.pri	[chemical] literature abstracts; [chemical] literature citations.
data.pri	[biological] literature abstracts; [biological] literature citations.	data.sec	[chemical] index terms.
data.sec	[biological] index terms.	gen.nam	CAS Customer Service
gen.nam	BIOSIS User Services	gen.add	Chemical Abstracts Service
gen.add	2100 Arch Street Philadelphia, PA 19103-1399 U.S.A.		P.O. Box 3012
gen.tel	215-587-4800		Columbus, Ohio 43210
///			U.S.A.
entry	BRD	gen.tel	800-848-6538
name.now	Berlin RNA Data Bank	///	
number	10002	entry	CASORF
		name.now	CAS ONLINE Registry File
		number	10051
		data.pri	[chemical] substances.
		data.sec	index terms; molecular formulae; [chemical] literature citations; molecular structure [chemical representations].
		gen.nam	CAS Customer Service
		gen.add	Chemical Abstracts Service
			P.O. Box 3012

	Columbus, OH 43210 U.S.A.	database characteristics; index terms; database maintenance [hardware]; database maintenance [software].
gen.tel	800-848-6538	Dr. Kathleen Arnett
///		Bioinformatics Dep't, A.T.C.C. 12301 Parklawn Drive Rockville, MD 20852 U.S.A.
entry	CCD	gen.net
name.now	Cambridge Structural Database	301-231-5527
number	10039	ka3@nihcu.bitnet
data.pri	[small molecule] atomic coordinates.	///
gen.nam	Dr. Olga Kennard	entry
gen.add	Cambridge Crystallographic Data Centre University Chemical Laboratory Lensfield Road Cambridge CB2 1EW U.K.	DCT
gen.tel	[011-44] 223-336409	name.now
gen.net	ok10%chemcrys@uk.ac.cam	Drosophila Codon Tables
///		number
entry	CGC	10041
name.now	Caenorhabditis Genetics Center	data.pri
number	10054	[Drosophila] codon usage; ['host' gene] nucleotide sequences; [transposable element ORF] nucleotide
data.pri	organisms [C. elegans strains].	sequences.
data.sec	[C. elegans] genetic maps; literature citations.	data.sec
gen.nam	Dr. Mark Edgley	[database] cross-references; ['host' gene] base composition;
gen.add	110 Tucker Hall University of Missouri Columbia, MO 65211	[transposable element ORF] base composition.
gen.tel	314-882-7384	gen.nam
gen.net	bioscgc@umcmb.bitnet	Dr. Michael Ashburner Department of Genetics University of Cambridge Cambridge U.K.
///		gen.tel
entry	CSRS	[011-44] 223-333-969
name.now	Compilation of Small RNA Sequences	gen.net
number	10004	ma11%cam.phx@uk.ac.cam.eng-ifc
data.pri	[small RNA] nucleotide sequences.	///
gen.nam	Dr. Ram Reddy	entry
gen.add	Department of Pharmacology Baylor College of Medicine Houston, TX 77030 U.S.A.	DDBJ
gen.tel	713-799-4458 ext. 26	name.now
///		DNA Data Bank of Japan
entry	CUTG	number
name.now	Codon Usage Tabulation from GENBANK	10006
number	10005	data.pri
data.pri	codon usage.	nucleotide sequences.
data.sec	[GENBANK] cross-references.	data.sec
gen.nam	Dr. Toshimichi Ikemura	[nucleotide sequence] functional features; index terms; [molecular biology]
gen.add	National Institute of Genetics Mishima, Shizuoka 411 JAPAN	literature citations; taxonomic identification.
gen.tel	[011-81] 559-75-0771, ext. 355	gen.nam
///		Dr. Sanzo Miyazawa National Institute of Genetics Mishima 411 JAPAN
entry	DBIR	gen.tel
name.now	Directory of Biotechnology Information/Resources	[011-81] 559-75-0771
number	10053	gen.net
data.pri	[biotechnology] databases.	ddb%nigsun.uucp@vax2.nlm.nih.gov
data.sec	database access information;	///
		entry
		DRHPL
		name.now
		Database for the Repository of Human Probes and Libraries
		number
		10007
		data.pri
		[human cDNA] clones; RFLPs; clones [chromosome specific libraries]; [human genomic] clones.
		data.sec
		clones [descriptions]; literature citations; clones [library characterizations]; clones [distribution lists].
		gen.nam
		Dr. Donna R. Maglott
		gen.add
		American Type Culture Collection 12301 Parklawn Drive

	Rockville, MD 20852-1776 U.S.A.	Box 3426 D-3400 Gottingen WEST GERMANY
gen.tel	301-231-5586	gen.tel [011-49] 551-397-599
///		///
entry	DROSO	entry GDN
name.now	Genetic Variations of Drosophila melanogaster	name.now Gene Diagnosis Newsletter
number	10033	number 10060
data.pri	[D. melanogaster] genetic variants.	data.pri [disease, human inherited] literature citations.
data.sec	[D. melanogaster] chromosome rearrangements; [D. melanogaster] genetic mutations; [D. melanogaster non-chromosomal] phenotypes;	data.sec — gen.nam Dr. Jorg Schmidtke gen.add Gene Diagnosis Newsletter
organism	[D. melanogaster, wild-type strains].	Box 3426 D-3400 Gottingen WEST GERMANY
gen.nam	Dr. Daniel L. Lindsley	gen.tel [011-49] 551-397-599
gen.add	Department of Biology University of California at San Diego La Jolla, CA 92093 U.S.A.	///
///		entry GENBANK
entry	EMBL	name.now The GenBank Genetic Sequence Data Bank
name.now	The EMBL Data Library	number 10009
number	10008	data.pri nucleotide sequences.
data.pri	nucleotide sequences.	data.sec [nucleotide sequence] functional features; index terms; [molecular biology]
data.sec	[nucleotide sequence] functional features; [molecular biology] literature citations; index terms; taxonomic identification.	literature citations; taxonomic identification.
gen.nam	EMBL Data Library	gen.nam Dr. David Benton
gen.add	Postfach 10.2209 6900 Heidelberg WEST GERMANY	gen.add IntelliGenetics, Inc. 700 El Camino Real East Mountain View, CA 94040 U.S.A.
gen.tel	[011-49] 6221-387-258	gen.tel 415-962-7364
gen.net	datalib@embl.bitnet	gen.net genbank@bionet-20.bio.net
///		///
entry	GBSOFT	entry HDB
name.now	The GenBank Software Clearinghouse	name.now Hybridoma Data Bank: A Data Bank on Immunoclonies
number	10028	number 10010
data.pri	literature citations [software].	data.pri hybridomas; clones [immuno-]; [monoclonal] antibodies.
data.sec	software [acquisition information]; software [system requirements]; software [functions].	data.sec index terms; literature citations.
gen.nam	GenBank Software Clearinghouse	gen.nam Dr. Esther J. Asaki
gen.add	IntelliGenetics, Inc. 700 El Camino Real East Mountain View, CA 94040 U.S.A.	gen.add Hybridoma Data Bank A.T.C.C. 12301 Parklawn Drive Rockville, MD 20852 U.S.A.
gen.tel	415-962-7364	gen.tel 301-231-5527
gen.net	genbank@bionet-20.bio.net	gen.net zja@nihcu.bitnet
///		///
entry	GC	entry HGIR
name.now	Gene Communications	name.now Human Genome Information Resource
number	10052	number 10058
data.pri	[nucleotide sequence, cloned] literature citations.	data.pri nucleotide probes; restriction fragments; physical maps [data].
data.sec	—	gen.nam Dr. Thomas Marr
gen.nam	Dr. Jorg Schmidtke	gen.add MS K710
gen.add	Gene Communications	

	Los Alamos National Laboratory Los Alamos, NM 87545	[molecular biology] literature citations; [amino acid mutagenizing] methods; [amino acid] chemical modification; [protein] molecular properties; [protein variant] sources.
gen.tel	505-665-0838	Dr. Akira Tsugita
gen.net	tgm@hedonist-1.lanl.gov	JIPID
///		Life Science Institute Science University of Tokyo Yamazaki Noda 278 JAPAN
entry	HGMCR	gen.nam
name.now	NIGMS Human Genetic Mutant Cell Repository Catalog of Cell Lines	gen.add
number	10034	gen.tel
data.pri	[human] cell lines.	[011-81] 471-24-1501 ext. 5001
data.sec	cell lines [characteristics]; index terms.	[011-81] 471-22-3899
gen.nam	Dr. Arthur E. Greene	gen.net
gen.add	Coriell Institute for Medical Research Copewood and Davis Streets Camden, NJ 08103 U.S.A.	ex5292@jpnst30.bitnet 42:cdt0079 (CODATA/Dialcom)
gen.tel	609-966-7377	///
///		entry JIPIDB
entry	HGML	name.now Biological Database: Asian and Oceania node of the International Protein Sequence Database
name.now	The Howard Hughes Medical Institute Human Gene Mapping Library	number 10044
number	10011	data.pri [protein] molecular properties.
data.pri	[human] DNA probes; [human] RFLPs; [molecular biology] literature citations; [human] genetic maps.	data.sec [database] cross-references; [biological] literature citations; [protein] sources; index terms.
data.sec	index terms.	gen.nam Dr. Akira Tsugita
gen.nam	Ms. Iva H. Cohen	gen.add JIPID
gen.add	Human Gene Mapping Library 25 Science Park New Haven, CT 06511 U.S.A.	Life Science Institute Science University of Tokyo Yamazaki Noda 278 JAPAN
gen.tel	203-786-5515	gen.tel [011-81] 471-24-1501 ext. 5001
gen.net	genesic@yalevm.bitnet	[011-81] 471-22-3899
///		gen.net ex5292@jpnst30.bitnet 42:cdt0079 (CODATA/Dialcom)
entry	HIVSSA	///
name.now	HIV Sequence & Sequence Analysis Database	entry JIPIDM
number	10040	name.now NMR Database on biopolymers: Asian and Oceania node of the International Protein Sequence Database
data.pri	[HIV] amino acid sequences; [HIV] nucleotide sequences; sequence alignments.	number 10045
data.sec	AIDS related animal viruses.	data.pri NMR data; literature citations.
gen.nam	Ms. Kersti MacInnes	data.sec [database] cross-references; index terms; NMR data interpretation.
gen.add	T-10, MS K710 Los Alamos National Laboratory Los Alamos, NM 87545 U.S.A.	gen.nam Dr. Akira Tsugita
gen.tel	505-667-7510	gen.add JIPID
gen.net	kam@lanl.gov	Life Science Institute Science University of Tokyo Yamazaki Noda 278 JAPAN
///		gen.tel [011-81] 471-24-1501 ext. 5001
entry	JIPIDA	[011-81] 471-22-3899
name.now	Artificial Variant Database: Asian and Oceania node of the International Protein Sequence Database	gen.net ex5292@jpnst30.bitnet 42:cdt0079 (CODATA/Dialcom)
number	10038	///
data.pri	amino acid sequences [artificial variants].	entry JIPIDN
data.sec	[database] cross-references; index terms;	name.now Natural Variant Database: Asian and

	Oceania node of the International Protein Sequence Database		Los Alamos, NM 87545
number	10046	U.S.A.	
data.pri	amino acid sequences [natural variants]. [database] cross-references; index terms; [molecular biology] literature citations; [amino acid] chemical modification; [protein variant] molecular properties; [protein variant] sources.	gen.tel	505-667-6683
data.sec		gen.net	limb@lanl.gov
		///	
gen.nam	Dr. Akira Tsugita	entry	LYSIS
gen.add	JIPID	name.now	Proteolysis Database
	Life Science Institute Science University of Tokyo Yamazaki Noda 278 JAPAN	number	10061
gen.tel	[011-81] 471-24-1501 ext. 5001 [011-81] 471-22-3899	data.pri	[protein substrate cleavage site] literature citations; proteinase specificity.
gen.net	ex5292@jpsut30.bitnet 42:cdt0079 (CODATA/Dialcom)	data.sec	peptides [synthetic substrates].
///		gen.nam	Dr. Borivoj Keil
entry	JIPIDS	gen.add	Institut Pasteur
name.now	Sequence Database: Asian and Oceania node of the International Protein Information Database		Chimie des Proteines
number	10047	gen.tel	25 rue du docteur Roux
data.pri	nucleotide sequences; amino acid sequences.	gen.net	Paris 75015
data.sec	[database] cross-references; index terms; [nucleotide sequence] functional features; [amino acid sequence] functional features; [biological] literature citations; [sequence] sources.		FRANCE
gen.nam	Dr. Akira Tsugita	gen.tel	[011-33] 1-4734-0593
gen.add	JIPID	gen.net	keil@pasteur.bitnet
	Life Science Institute Science University of Tokyo Yamazaki Noda 278 JAPAN		dbi0061.dialcom (Telecom Gold)
gen.tel	[011-81] 471-24-1501 ext. 5001 [011-81] 471-22-3899	///	
gen.net	ex5292@jpsut30.bitnet 42:cdt0079 (CODATA/Dialcom)	entry	MEDLINE
///		name.now	MEDLINE and Backfiles
entry	LIMB	number	10013
name.now	LiMB	ata.pri	[medical] index terms; [medical] literature citations;
number	10012	data.sec	[biological] index terms; [biological] literature citations.
data.pri	[molecular biology] databases.	gen.nam	[medical] literature abstracts; [biological] literature abstracts.
data.sec	database access information; database characteristics; [molecular biology] literature citations; database maintenance [hardware]; database maintenance [software]; database contribution information.	gen.add	MEDLARS Management Section
gen.nam	Dr. Christian Burks		National Library of Medicine
gen.add	Theoretical Biology and Biophysics T-10, MS K710 Los Alamos National Laboratory	gen.tel	Bldg. 38A
			National Institutes of Health
			Bethesda, MD 20894
			U.S.A.
		gen.tel	800-638-8480
		///	
entry	MIM	entry	MIM
name.now	Mendelian Inheritance in Man	name.now	Mendelian Inheritance in Man
number	10064	number	10027
data.pri	[human inherited] diseases.	data.pri	[cultured] microorganisms.
gen.nam	Dr. Victor McKusick	data.sec	taxonomic classification; [microorganism] sources; microorganism [history]; literature citations.
gen.add	Moore Clinic		
		gen.tel	301-955-6641
		///	
entry	MICIS	entry	MICIS
name.now	Microbial Culture Information Service	name.now	Microbial Culture Information Service
number	10027	number	10027
data.pri	[cultured] microorganisms.	data.pri	[cultured] microorganisms.
data.sec	taxonomic classification; [microorganism] sources; microorganism [history]; literature citations.	data.sec	taxonomic classification; [microorganism] sources; microorganism [history]; literature citations.

gen.nam	Dr. Geraldine Alliston	number	10014
gen.add	Laboratory of the Government Chemist	data.pri	amino acid sequences.
	Cornwall House	data.sec	[molecular biology] literature citations;
	Waterloo Road		[amino acid sequence] sources.
	London SE1 8XY	gen.nam	Dr. R.F. Doolittle
	U.K.	gen.add	Chemistry Department D-006
gen.tel	[011-44] 1-211-8834		University of California, San Diego
///			La Jolla, CA 92093
entry	MIPS	U.S.A.	
name.now	Martinsreid Institute for Protein Sequence Data	gen.tel	619-534-4417
number	10037	///	
data.pri	amino acid sequences.	entry	PDB
data.sec	[amino acid sequence] literature citations; [amino acid sequence] functional features; index terms.	name.now	Protein Data Bank
gen.nam	Dr. Hans-Werner Mewes	number	10015
gen.add	Max Planck Institut fur Biochemie	data.pri	[biomacromolecular] atomic coordinates.
	MPI/GEN	data.sec	[biomacromolecule] functional features; [biomacromolecular] literature citations; [biomacromolecule] molecular properties.
	8033 Martinsreid	gen.nam	Dr. Frances C. Bernstein
	WEST GERMANY	gen.add	Chemistry Department
gen.net	mewes@dm0mpb51.bitnet		Brookhaven National Laboratory
///			Upton, NY 11973
entry	MSDN	gen.tel	U.S.A.
name.now	Microbial Strain Data Network	gen.net	516-282-4382
number	10032	///	abola@bnldag.bitnet
data.pri	[microbial strain database] cross-references; [microbial strain] index terms.	entry	PIR
data.sec	database access information; database characteristics; database contribution information.	name.now	National Biomedical Research Foundation
gen.nam	MSDN Secretariat		Protein Identification Resource
gen.add	Biotechnology Centre		Protein Sequence Database
	Cambridge University	number	10016
	307 Huntingdon Rd	data.pri	amino acid sequences.
	Girton, Cambridge CB3 0JX	data.sec	[amino acid sequence] literature citations; [amino acid sequence] functional features; index terms.
	U.K.	gen.nam	Dr. Kathryn E. Sidman
gen.tel	[011-44] 223-277-502	gen.add	National Biomedical Research Foundation
gen.net	42:cdt0001 (CODATA/Dialcom)		3900 Reservoir Road, N.W.
///			Washington, D.C. 20007
entry	NAPRALERT	gen.tel	U.S.A.
name.now	NAPRALERT	gen.net	202-687-2121
number	10059	///	pirmail@gunbrf.bitnet
data.pri	[chemical compound] literature citations.	entry	PPR
data.sec	[taxonomy] literature citations; [pharmacology] literature citations.	name.now	Plasmid Prefix Registry
gen.nam	Dr. Norman Farnsworth	number	10017
gen.add	University of Illinois at Chicago	data.pri	plasmid prefixes.
	Prog. for Collaborative Research in the Pharmaceutical Sciences	data.sec	plasmid prefixes [locale of registrants]; plasmid prefixes [name of registrants].
	833 S. Wood St.	gen.nam	Dr. Esther Lederberg
	Chicago, IL 60612	gen.add	Department of Medical Microbiology
gen.tel	312-996-7253		Sherman Fairchild Building - 5402
gen.net	beecher@uicbal.bitnet		Stanford University School of Medicine
///			Stanford, CA 94305-2499
entry	NEWAT	gen.tel	U.S.A.
name.now	NEWAT	///	415-723-1772

entry	<b>PRFLITDB</b>	gel electrophoretic data
name.now	<b>PRF/LITDB</b>	[protein, spot quality];
data.pri	[amino acid sequence] literature abstracts;	[protein] molecular properties.
	[amino acid sequence] literature citations.	
data.sec	index terms.	
gen.nam	Dr. Yasuhiko Seto	Dr. Cecile Chang
gen.add	Protein Research Foundation	Cold Spring Harbor Laboratory
	4-1-2 Ina, Minoh-shi	P.O. Box 100
	Osaka 562	Cold Spring Harbor, NY 11724
	JAPAN	U.S.A.
///		516-367-8356
entry	<b>PRFSEQDB</b>	///
name.now	<b>PRF/SEQDB</b>	entry RED
number	10018	name.now Restriction Enzyme Database
data.pri	amino acid sequences.	number 10022
data.sec	[amino acid sequence] functional features;	data.pri restriction enzymes.
	[biological] literature citations.	data.sec [restriction enzyme] functional features;
gen.nam	Dr. Yasuhiko Seto	literature citations; literature abstracts;
gen.add	Protein Research Foundation	[restriction enzyme isolation] methods.
	4-1-2 Ina, Minoh-shi	gen.nam Dr. Rich Roberts
	Osaka 562	gen.add Cold Spring Harbor Laboratory
	JAPAN	Cold Spring Harbor, NY 11724
///		U.S.A.
entry	<b>PSEQIP</b>	gen.tel 516-367-8388
name.now	PseqIP	gen.net roberts@bionet-20.arpa
number	10019	///
data.pri	amino acid sequences.	entry RFLPD
data.sec	index terms.	name.now RFLP Database
gen.nam	Dr. Isabelle Sauvaget	number 10057
gen.add	Institut Pasteur	data.pri RFLPs.
	25 rue du Docteur Roux	gen.nam Dr. Howard Cann
	75724 Paris Cedex 15	gen.add Centre d'Etude du Polymorphisme
	FRANCE	Humain
gen.tel	[011-33] 1-45-68-85-09	3, rue d'Ulm
///		75005 Paris
entry	<b>PTG</b>	FRANCE
name.now	Protein Translation of GENBANK	gen.tel [011-33] 1-43-29-12-11
number	10020	///
data.pri	amino acid sequences.	entry SEQANALREF
data.sec	[nucleotide sequence] functional features;	name.now Sequence Analysis Literature Reference
	[molecular biology] literature citations;	Data Bank
	index terms; taxonomic identification.	number 10055
gen.nam	Dr. Christian Burks	data.pri [sequence analysis] literature citations.
gen.add	Theoretical Biology and Biophysics Grp.	data.sec index terms.
	T-10, MS K710	gen.nam Dr. Amos Bairoch
	Los Alamos National Laboratory	gen.add Departement de Biochimie Medicale
	Los Alamos, NM 87545	C.M.U.
	U.S.A.	1 Rue Michel Servet
gen.tel	505-667-6683	1211 Geneva 4
gen.net	cb@lanl.gov	SWITZERLAND
///		gen.tel [011-41] 22-61-84-92
entry	<b>QTDPGD</b>	gen.net bairoch@cgecmu51.bitnet
name.now	Quest 2D Gel Protein Database	///
number	10021	entry SIGPEP
data.pri	gel electrophoretic data	name.now SIGPEP
	[protein, spot representation];	number 10029
data.sec	gel electrophoretic data	data.pri [signal peptide] amino acid sequences.
	[protein, spot coordinates];	data.sec [amino acid sequence] sources;
		literature citations.
		gen.nam Dr. Gunnar von Heijne

gen.add	Theoretical Physics Royal Institute of Technology S-10044 Stockholm SWEDEN	gen.nam	properties; [chemical synthesis] methods.
gen.tel	[011-46] 8-787 7172	gen.add	Dr. Philip Wexler
///			National Library of Medicine
entry	SRRSD		8600 Rockville Pike
name.now	16S Ribosomal RNA Sequence Database		Bethesda, MD 20894
number	10024		U.S.A.
data.pri	[16S rRNA] nucleotide sequences.	gen.tel	301-496-6531
data.sec	[16S rRNA nucleotide sequence] functional features.	///	
gen.nam	Dr. George Fox	entry	TRF
gen.add	Department of Biochemical Science University of Houston SR-1 4800 Calhoun Houston, TX 77004 U.S.A.	name.now	The Taxonomic Reference File at BIOSIS
gen.tel	713-749-3980; 713-749-2830	number	10048
gen.net	fox@uhrc2.bitnet	data.pri	organism [bacterial].
	fox@uhrc2.crcc.uh.edu	data.sec	organism [bacterial, synonyms]; taxonomic classification; index terms.
///		gen.nam	Dr. Robert T. Howey
entry	SRSRSC	gen.add	BIOSIS
name.now	Small Ribosomal Subunit RNA Sequence Compilation		2100 Arch Street
number	10025		Philadelphia, PA 19103-1399
data.pri	[small subunit rRNA] nucleotide sequences.	gen.tel	U.S.A.
data.sec	[nucleotide sequence, small subunit rRNA] functional features; [sequence] sources; [molecular biology] literature citations.	gen.net	215-587-4917
gen.nam	Dr. Rupert De Wachter		42:CDT0018 (CODATA/Dialcom)
gen.add	Departement Biochemie Universiteit Antwerpen (UIA) Universiteitsplein 1 B-2610 Antwerpen BELGIUM	///	
gen.tel	[011-32] 3-828-25-28, ext. 269	entry	TRNAC
///		name.now	tRNA Compilation
entry	SWISSPROT	number	10026
name.now	SWISS-PROT Protein Sequence Data Bank	data.pri	[tRNA] nucleotide sequences; [tRNA gene] nucleotide sequences; sequence alignments.
number	10023	data.sec	[molecular biology] literature citations; [tRNA] secondary structures; [tRNA] sources.
data.pri	amino acid sequences.	gen.nam	Dr. Mathias Sprinzl
data.sec	[amino acid sequence] functional features; index terms.	gen.add	Department of Biochemistry University of Bayreuth Universitätsstrasse 30
gen.nam	Dr. Amos Bairoch		D-8580 Bayreuth
gen.add	Departement de Biochimie Medicale C.M.U. 1 Rue Michel Servet 1211 Geneva 4 SWITZERLAND	gen.tel	WEST GERMANY
gen.tel	[011-41] 22-61-84-92	gen.net	[011-49] 921-552-668
gen.net	bairoch@cgecmu51.bitnet		datex-p 45921040258
///		///	
entry	TOXNET	entry	VECTOR
name.now	Toxicology Data Network	name.now	Cloning Vector Sequence Database
number	10049	number	10036
data.pri	[toxic] chemicals.	data.pri	[cloning vector] nucleotide sequences.
data.sec	[chemical] literature citations; toxin	data.sec	[molecular biology] literature citations; [cloning vector nucleotide sequence] base composition; [cloning vector] sources; [database] cross-references; index terms.
gen.nam	Dr. William Gilbert	gen.nam	Dr. William Gilbert
gen.add	Whitehead Institute, Room 211 Nine Cambridge Center Cambridge, MA 02142 U.S.A.	gen.add	Whitehead Institute, Room 211 Nine Cambridge Center Cambridge, MA 02142
gen.tel	617-258-5139		U.S.A.
gen.net	gilbert@mitwibr.bitnet		
///			

## Nucleic Acids Research

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taxonomic classification  
taxonomic identification  
toxin properties

... MICIS, TRF  
... DDBJ, EMBL, GENBANK, PTG  
... TOXNET

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