Protein Data Bank

Quarterly Newsletter

Number 62

October 1992

October Update

The Protein Data Bank (PDB) is pleased to announce the availability of atomic coordinate entries on CD ROM, and on-line from Brookhaven National Laboratory (BNL) by anonymous FTP and the PDB e-mail file server.

The latest October 1992 PDB release includes over 1,000 fully annotated coordinate entries together with more than 300 pre-release entries (see below). Both the fully annotated and pre-release coordinate entries are now accessible on-line from BNL. The PDB provides this on-line service free of charge. It is intended to meet the needs of users requiring access to a limited number of data entries. Individuals needing the complete release should continue to order the PDB on magnetic tape or on the newly available PDB CD ROM. Besides the atomic coordinates, the CD ROM includes the 290 available structure factor entries in compressed format.

The PDB Newsletter contains a Brookhaven Order Form with current prices that may be used to place an order for data on CD or tape. It is also possible to obtain copies of the Order Form in electronic form on-line from BNL, or in hard copy by contacting the PDB at the address given.

October 1992 PDB Release

1007 fully annotated atomic coordinate entries (53 new atomic coordinate entries)

897 proteins, enzymes and viruses

- 89 DNA's
- 2 RNA's
- 9 tRNA's
- 10 carbohydrates

303 pre-release atomic coordinate entries

- 290 structure factor entries
- 23 NMR experimental entries

Atomic Coordinate Entries in Preparation

- 111 entries awaiting approval
- 43 entries on hold

To Contact the PDB

Protein Data Bank Chemistry Department, Building 555 Brookhaven National Laboratory Upton, NY 11973 USA

Telephone: +1 516-282-3629 Facsimile: +1 516-282-5751 e-mail: pdb@bnlchm.bitnet *or* pdb@chm.chm.bnl.gov

Please include your telephone number, facsimile number, mailing address, and e-mail address in all correspondence.

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The tables on pp. 3-4 of this Newsletter are abbreviated and contain only new information. Full versions of all PDB tables are available from the PDB anonymous FTP and e-mail server or by normal mail upon request. Please see the Brookhaven Order Form on pp. 5-6 for requesting various PDB documentation.

PDB Staff News

We are pleased to announce that Nancy Oeder has joined the PDB staff as a member of the database production group. Nancy comes to us from Rensselaer Polytechnic Institute where she worked on the structure of gramicidin-A under the direction of Professor Bonnie Wallace. Nancy's appointment to the staff will help us in our work to ensure that the PDB remains current and that IDENT Codes are issued promptly for newly deposited entries.

Accessing PDB via Gopher

The PDB anonymous FTP directories now can be accessed via the Gopher service. The PDB Internet Gopher server allows access to coordinate entries through a window environment. While it is possible to identify and acquire a desired file by browsing through the directories, a full-index search will be on-line shortly. The Internet Gopher is a distributed document delivery service. It allows the novice user to access various types of data residing on multiple hosts. This service uses a client-server model in which users are presented with a hierarchical representation of directories of documents. The user is the "client" who makes queries to a "server" residing on another host on the Internet network. This system has been set up so that the user does not need to know the location of servers, since Gopher transparently handles the connections.

Information for setting up an Internet Gopher client including source files for different machines is available from the PDB anonymous FTP server. Those now running a Gopher client can access the PDB FTP site by including the following link:

Name = Protein Data Bank FTP site

Type = 1

Host = pdb.pdb.bnl.gov

Port = 70

Path = 1/

For more information or help in accessing the PDB via Gopher, you may send an e-mail message to oeder@bnl.gov.

File Server and Anonymous FTP

The PDB e-mail file server and anonymous FTP are available for all users. In addition to the atomic coordinate entry files, it is possible to access and download PDB general information and documentation. For more information about the e-mail server, send a message to:

fileserv@pb1.pdb.bnl.gov

and include the following text:

'send info your_e-mail_address'.

The PDB also has an anonymous FTP account available on the system pdb.pdb.bnl.gov with Internet address 130.199.144.1. Files may be transferred to and from this system using 'anonymous' as the FTP user name and your real user name as the password. Besides downloading data files and documentation, it is possible to upload any files you may wish to send to the PDB. Please note that those using VMS may need to place quotes around file names.

Anyone experiencing problems or having questions related to the above network services may send an email message to skora@bnl.gov.

CD ROM Information

The October release and all future releases will be available on CD ROM in ISO 9660 format. To avoid confusion, the layout of the files on the CD ROM will mirror that of the tape distribution. The entry files themselves are in ASCII format and should be readable by any software able to read text files. The structure factor files will be compressed using the standard UNIX compress command.

There is currently no software provided on the CD ROM for accessing the data files, although we expect this to change in the future.

VAX/VMS systems currently do not directly support access to ISO 9660 formatted CD ROMs. The PDB CD ROM may be accessed on VAX/VMS systems by two approaches:

- 1. There is an ISO 9660 compliant device driver available from Digital Equipment Corporation (DEC) that allows direct access to the CD ROM (driver part number YT-GS001-01). Please contact your DEC sales representative for further information.
- 2. There is a public utility for accessing ISO 9660 CD ROMs, called CD_ACCESS, written by Peter Stockwell, University of Otago, New Zealand, that will allow all the files on the CD ROM to be copied to a magnetic disk drive. This utility can be obtained from the EMBL e-mail server (for further information you may contact DataLib@EMBL-Heidelberg.DE). When copying files using CD_ACCESS, be sure to use the /BINARY qualifier to the copy command.

Newly Released Entries

1LPE	APOLIPOPROTEIN E3(LDL RECEPTR-BNDNG DMN)	C WILCON D ACARD
1LE2	APOLIPOPROTEIN E2(LDL RECEPTR-BNDNG DMN)	C.WILSON,D.AGARD
1LE4	APOLIPOPROTEIN E4(LDL RECEPTR-BNDNG DMN)	C.WILSON,D.AGARD
5ABP	L-ARABINOSE-BINDING PROTEIN/D-GALACTOSE	C.WILSON,D.AGARD
4CA2	CARBONIC ANHYDRASE II(HUMAN RECOMBINANT)	F.QUIOCHO,D.WILSON,N.VYAS
5CA2	CARBONIC ANHYDRASE II MUTANT(T200S)	R.ALEXANDER, D. CHRISTIANSON
12CA	CARBONIC ANHYDRASE II MUTANT(12003)	R.ALEXANDER, D.CHRISTIANSON
1CA3	CARBONIC ANHYDRASE II MOTANT(V12TA) CARBONIC ANHYDRASE II (HUMAN)(PH 5.7)	S.NAIR,D.CHRISTIANSON
1HCA	, , ,	S.NAIR,D.CHRISTIANSON
1HEA	CARBONIC ANHYDRASE II(HUMAN)(PH 6.5)	S.K.NAIR,D.W.CHRISTIANSON
1HEB	CARBONIC ANHYDRASE II MUTANT(L198R)	S.K.NAIR,D.W.CHRISTIANSON
1HEC	CARBONIC ANHYDRASE II MUTANT(L198E)	S.K.NAIR,D.W.CHRISTIANSON
1HED	CARBONIC ANHYDRASE II MUTANT(L198H)	S.K.NAIR,D.W.CHRISTIANSON
	CARBONIC ANHYDRASE II MUTANT(1L98A)	S.K.NAIR,D.W.CHRISTIANSON
3CMS	CHYMOSIN B MUTANT(V111F)(BOVINE)	T.BLUNDELL ET AL.
1CGC	DNA(B,CCGGCGCGG,SYNTHETIC)	U.HEINEMANN,M.BANSAL
1D58	DNA(TGATCA,SYNTH) HEPIADRIAMYCIN	B.L.D'ESTAINTOT,B.GALLOIS,T.BROWN,W.N.HUNTER
1D63	DNA(CGCAAATTTGCG,SYNTH)/BERENIL	D.BROWN,S.NEIDLE ET AL.
1D64	DNA(CGCGAATTCGCG,SYNTH)/PENTAMIDINE	EDWARDS, JENKINS, NEIDLE
1D65	DNA(CGCAAATTTGCG,SYNTHETIC)	K.EDWARDS,S.NEIDLE ET AL.
1D81	DNA(CGCAAATTIGCG,SYNTH): I=INOSINE	G.A.LEONARD,E.D.BOOTH,W.N.HUNTER,T.BROWN
1D82	DNA(GTCTAGAC,SYNTHETIC)	A.CERVI,B.LANGOLOIS D'ESTAINTOT,W.HUNTER
10FX	RNA(GCG)D(TATACCC)/D(GGGTATACGC)OKAZAKI	EGLI,USMAN,ZHANG,RICH
8EST	ELASTASE(PORCINE)/GUANIDINIUM ISOCOUMARIN	R.RADHAKRISHNAN,E.MEYER JR
1GPA	GLYCOGEN PHOSPHORYLASE A(R STATE)	BARFORD,HU,JOHNSON
1GPB	GLYCOGEN PHOSPHORYLASE B	JOHNSON,ACHARYA,STUART
2GPB	GLYCOGEN PHOSPHORYLASE B/GLC	J.MARTIN,L.JOHNSON
3GPB	GLYCOGEN PHOSPHORYLASE B/G1P	J.MARTIN,L.JOHNSON
4GPB	GLYCOGEN PHOSPHORYLASE B/GFP	J.MARTIN,L.JOHNSON
5GPB	GLYCOGEN PHOSPHORYLASE B/GMP/GLC	J.MARTIN,L.JOHNSON
6GPB	GLYCOGEN PHOSPHORYLASE B/H2P	L.JOHNSON,K.ACHARYA
7GPB	GLYCOGEN PHOSPHORYLASE B(R STATE)/AMP	BARFORD,HU,JOHNSON
8GPB	GLYCOGEN PHOSPHORYLASE B(T STATE)/AMP	BARFORD,HU,JOHNSON
9GPB	GLYCOGEN PHOSPHORYLASE B(R STATE)	BARFORD, JOHNSON
1NIH	HEMOGLOBIN(ALPHA-NICKEL,BETA-FERROUS)	B.LUISI,B.LIDDINGTON
1PBX	HEMOGLOBIN(PAGOTHENIE BERNACCHII,CARBOMONOXY)	G.FERMI
1HSA	HISTOCOMPATIBILITY ANTIGEN(HUMAN)/HLA-B-2705	D.R.MADDEN, J.C.GORGA, J.L.STROMINGER, D.C.WILEY
1HHP	HIV-1 PROTEASE(ISOLATE BRU)	S.SPINELLI,P.M.ALZARI
1FAI	IGG2B FAB(KAPPA) FROM A MONOCLONAL ANTI-ARSONATE AB	M.B.LASCOMBE, P.M.ALZARI, R.J. POLJAK, A.NISONOFF
	(STRUCTURE 1)	
2F19	IGG2B FAB(KAPPA) FROM A MONOCLONAL ANTI-ARSONATE AB	M.B.LASCOMBE,P.M.ALZARI,R.J.POLJAK,A.NISONOFF
011 A	(STRUCTURE 2)	
2ILA 6I1B	INTERLEUKIN 1A(HUMAN)	B.GRAVES,M.HATADA
711B	INTERLEUKIN 1B(HUMAN)(NMR,AVERAGED STRC)	CLORE, WINGFIELD, GRONENBORN
	INTERLEUKIN 1B(HUMAN)(NMR,32 STRUCTURES)	CLORE, WINGFIELD, GRONENBORN
3IL8	INTERLEUKIN 8(HUMAN)	A.WLODAWER
2ZTA	LEUCINE ZIPPER(GCN4 TAP)	O'SHEA,KLEMM,KIM,ALBER
1MEE	MESENTERICOPEPTIDASE/EGLIN C	Z.DAUTER,C.BETZEL,K.WILSON
2PSG	PEPSINOGEN(PORCINE)	M.JAMES,A.SIELECKI
3RUB	RUBISCO(FORM III)	EISENBERG,SCHREUDER ET AL.
4RUB	RUBISCO(FORM IV)	EISENBERG,SCHREUDER ET AL.
1STP	STREPTAVIDIN-BIOTIN COMPLEX	DUPONT PROTEIN CRYSTALLOGRAPHY
4TIM	TIM(TRYPANOSOMA)/2-PHOSPHOGLYCERATE	NOBLE, WIERENGA, HOL ET AL.
5TIM	TIM(TRYPANOSOMA)/SULFATE	R.WIERENGA,W.HOL ET AL.
6TIM	TIM(TRYPANOSOMA)/GLYCEROL-3-PHOSPHATE	NOBLE, WIERENGA, HOL ET AL.
2APD	APOLIPOPROTEIN D (MODEL)	M.C.PEITSCH,M.S.BOGUSKI

Newly Received Depositions

2AT2	ASPARTATE TRANSCARBAMOYLASE	1KST	KISTRIN (NMR)
1BIA	BIOTIN REPRESSOR	9LDB	LACTATE DEHYDROGENASE/NADH
1BIB	BIOTIN REPRESSOR/BIOTINYLATED LYSINE COMPLEX	9LDT	LACTATE DEHYDROGENASE/NADH/OXAMATE
1APO	BLOOD COAGULATION FACTOR X(BOVINE)N-TERMINAL EGF-LIKE MODULE(NMR, 13 STRUCTURES)	1LLD	OXIDOREDUCTASE(CHOH (D)-NAD (A))L-LACTATE DEHYDROGENASE MUTANT(C2105)/NADH
2BBM	CALMODULIN/CALMODULIN-BINDING DMN OF RABBIT SKELETAL MYOSIN LT CHAIN KINASE(MMR)	1LAA	LYSOZYME(HUMAN) MUTANT (D53E)
1CAN		1TAY	LYSOZYME(HUMAN) MUTANT(Y63A)
1CAN	CARBONIC ANHYDRASE(HUMAN) (REFINED BY TNT)	1TBY	LYSOZYME(HUMAN) MUTANT(Y63L)
1CAO	CARBONIC ANHYDRASE(HUMAN) (REFINED BY PROFFT)	1TCY	LYSOZYME(HUMAN) MUTANT(Y63F)
1CAH	CARBONIC ANHYDRASE(HUMAN)/BICARBONATE	1TDY	LYSOZYME(HUMAN) MUTANT(Y63W)
1CPS	CARBOXYPEPTIDASE A/SULFODIIMINE INHIBITOR	1L00	LYSOZYME(T4) MUTANT(Q105A)
3CD4	CD4(HUMAN) (RESIDUES 1-182)	1L98	LYSOZYME(T4) MUTANT(Q105E)
3GCR	GAMMA-IIIB CRYSTALLIN(BOVINE)	1L99	LYSOZYME(T4) MUTANT(Q105G)
2CPL	CYCLOPHILIN(HUMAN T CELL)	1LYE	LYSOZYME(T4) MUTANT(C54T,T59V,C97A)
1PHA	CYTOCHROME P450CAM(PSEUDOMONAS PUTIDA) CAMPHOR MONOXYGENASE/CAMPHOR(PLUS ISOMER)	1LYF	LYSOZYME(T4) MUTANT(C54T,T59S,C97A)
1PHB	CYTOCHROME P450CAM(PSEUDOMONAS PUTIDA)CAMPHOR	1LYG	LYSOZYME(T4) MUTANT(C54T,T59N,C97A)
	MONOXYGENASE/CAMPHOR(MINUS ISOMER)	1LYH	LYSOZYME(T4) MUTANT(C54T,T59G,C97A)
1PHC	CYTOCHROME P450CAM(PSEUDOMONAS PUTIDA) CAMPHOR	1LYI	LYSOZYME(T4) MUTANT(C54T,T59D,C97A)
	MONOXYGENASE	1LYJ	LYSOZYME(T4) MUTANT(C54T,T59A,C97A)
1PHD	CYTOCHROME P450CAM(P.PUTIDA) CAMPHOR MONOXYGENASE/2-PHENYL IMIDAZOLE(STRUCT 1)	2MSB	MANNOSE-BINDING PROTEIN A, LECTIN DOMAIN/CALCIUM AND MAN GLCNAC2ASN GLYCOPEPTIDE
1PHE	CYTOCHROME P450CAM(P.PUTIDA) CAMPHOR	2AAH	METHANOL DEHYDROGENASE(METHYLOPHILIS W3A1)
	MONOXYGENASE/2-PHENYL IMIDAZOLE(STRUCT 2)	1BBK	METHYLAMINE DEHYDROGENASE(PARACOCCUS DENITRIFICANS)
1PHF	CYTOCHROME P450CAM(PSEUDOMONAS PUTIDA) CAMPHOR MONOXYGENASE/4-PHENYL IMIDAZOLE	1MTA	METHYLAMINE DEHYDROGENASE/AMICYANIN/CYTOCHROME C551I
1PHG	CYTOCHROME P450CAM(PSEUDOMONAS PUTIDA) CAMPHOR	1SWM	MYOGLOBIN(SPERM WHALE, FERRIC)/AZIDE
iriid	MONOXYGENASE/METYAPONE	1PPN	PAPAIN CYS-25 WITH BOUND ATOM
1D67	DNA(TGATCA)/IDARUBICIN	1POL	BETA SUBUNIT OF POL III
1D80	DNA(CGCGAATTGGCG)	2PF1	PROTHROMBIN FRAGMENT 1(RESIDUES 1-156)
1D85	DNA(CGC(O6-ETHYL-G)AATTCGCG)/NETROPSIN (MG++ FORM)	1BRQ	APO RETINOL BINDING PROTEIN
1D86	DNA(CGCGAATTCGCG)/NETROPSIN COMPLEX (MG ⁺⁺ FORM)	1BRP	HOLO RETINOL BINDING PROTEIN
1D87	DNA-RNA CHIMERIC DUPLEX [R(G)D(CGTATACGC)]2 (RNA SYNTHETIC)	2RN2	RIBONUCLEASE H(ESCHERICHIA COLI)
1D88	DNA-RNA CHIMERIC DUPLEX [D(GCGT)R(A)D(TACGC)] (RNA SYNTHETIC)	1RGK	RIBONUCLEASE T1 MUTANT(E46Q)/2'-AMP
1EZM	ELASTASE ZINC METALLOPROTEASE	2AAD	RIBONUCLEASE T1 MUTANT(H40K)/GUANYLIC ACID
1END	T4 ENDONUCLEASE V	1APG	RICIN A CHAIN (CASTOR PLANT)/ADENYL(3'>5')GUANOSINE
1EED	ENDOTHIAPEPSIN/PD125754	1FMP	RICIN A CHAIN (CASTOR PLANT)/FORMYCIN 5'-MONOPHOSPHATE
1LTS	HEAT LABILE ENTEROTOXIN (LT); CHOLERA-LIKE TOXIN, AB5 TOXIN	1RBA	RUBISCO (RIBULOSE-1,5-BISPHOSPHATE
1LTT	HEAT LABILE ENTEROTOXIN (LT); CHOLERA-LIKE TOXIN, AB5 TOXIN/LACTOSE		CARBOXYLASE/OXYGENASE) MUTANT(D193N)
1FLV	FLAVODOXIN(ANABAENA 7120)	1SAS	SARCOPLASMIC CALCIUM-BINDING PROTEIN (ISOTYPE II)
10FV	FLAVODOXIN(ANACYSTIS NIDULANS,OXIDIZED)	1PTS	STREPTAVIDIN/PEPTIDE (FSHPQNT)
1FBA	FRUCTOSE-1,6-BISPHOSPHATE ALDOLASE	1SUB	SUBTILISIN BPN'CRB-S3 MUTANT(N218S,S221C)
1GRD	GLUCOCORTICOID RECEPTOR DNA-BINDING DOMAIN	1SUC	SUBTILISIN BPN'CRB-S3 MUTANT(N218S,S221C,M50F,Y217K)
1AAZ	GLUTAREDOXIN	1SUD	SUBTILISIN BPN'CRB-S3 MUTANT(M50F,Y217K,N218S,S221C)
1ABA	GLUTAREDOXIN MUTANT(V15G,Y16P)	1BAL	DIHYDROLIPOAMIDE SUCCINYLTRANSFERASE, E3-BINDING DOMAIN (NMR)
1GRC	GLYCINAMIDE RIBONUCLEOTIDE TRANSFORMYLASE	1TLK	TELOKIN(TURKEY)
1PYG	GLYCOGEN PHOSPHORYLASE (PYRIDOXAL-5'-PYROPHOSPHORYL DERIV.)	1ETR	E-THROMBIN(BOVINE)/MQPA
1HAM	HEMOGLOBIN(HUMAN) AALBORG	1ETS	E-THROMBIN(BOVINE)/NAPAP
1DXT	HEMOGLOBIN(HUMAN,DEOXY) (EXTRA N-TERM MET)	1ETT	E-THROMBIN(BOVINE)/TAPAP
1DXV	HEMOGLOBIN(HUMAN,DEOXY) MUTANT(V1A)	1TGI	TRANSFORMING GROWTH FACTOR BETA 2
	HEMOGLOBIN(HUMAN,DEOXY) MUTANT(VTA)	1BPU	TRYPSIN INHIBITOR(BOVINE,PANCREAS) MUTANT(N43G)
1DXU		1BBI	TRYPSIN/CHYMOTRYPSIN BOWMAN-BIRK INHIBITOR(NMR,AVERAGE)
1MHA	HISTOCOMPATIBILITY ANTIGEN (H-2K(B))(MOUSE CLASS I)	2BBI	TRYPSIN/CHYMOTRYPSIN BOWMAN-BIRK INHIBITOR(NMR,16 STRUCTURES)
1AAF	HIV-1 NUCLEOCAPSID PROTEIN, MN STRAIN	1TYA	TYROSYL TRNA SYNTHETASE MUTANT(T51A)
1BBO	HUMAN ENHANCER BINDING PROTEIN MUTANT(CIIABU)(NMR,60 STRUCTURES)	1TYB	TYROSYL TRNA SYNTHETASE MUTANT(T51G)
1IGM	IMMUNOGLOBULIN M(HUMAN) FV FRAGMENT	1TYC	TYROSYL TRNA SYNTHETASE MUTANT(T51P)
1HIL	IGG2A FAB FRAGMENT (FAB 17/9)	1TYD	TYROSYL TRNA SYNTHETASE MUTANT(T51S)
1HIM	IGG2A FAB FRAGMENT (FAB 17/9)	1UDP	URIDINE DIPHOSPHOGALACTOSE-4-EPIMERASE
1HIN	IGG2A FAB FRAGMENT (FAB 17/9)	1VAB	MHC CLASS I H-2K\UB\D AND SENDAI VIRUS
	INSULIN(HUMAN)(NMR)	1VAB	MHC CLASS I H-2K/UB/D AND VESICULAR STOMATITIS VIRUS
1HIU		2SNV	SINDBIS VIRUS CAPSID PROTEIN
1HIS	INSULIN(HUMAN) DES-PENTAPEPTIDE(NMR)	1CDA	MEMBRANE-BOUND GYLCOPROTEIN CD40/LIGAND MODEL
1HIT	INSULIN(HUMAN) MUTANT(F24G)(NMR)	1ITA	INTERLEUKIN-1 ALPHA(HUMAN) MODEL
1BBN	INTERLEUKIN 4(HUMAN)(NMR)		PROTEIN INHIBITOR C MODEL 2
		2PAI	FRO LEIN INFIDITOR O MODEL 2

	BROOKHAVEN ORDER FORM		
Name of User		Date	
Address		Phone	: .
		E-mail	
· · · · · · · · · · · · · · · · · · ·		Fax #	
DATAPRTP (all available coordinate entries, bibliographic entries, and some computer programs)	6250cpi 1600cpi TK50 VAX/VMS backup []\$492 []\$955 []\$653 VAX/VMS copy []\$492 []\$955 []\$653 Unlabeled ASCII []\$492 []\$955 Unlabeled EBCDIC []\$492 []\$955 SGI/SUN/IBM/E&S UNIX tar []\$223		<u>4mm</u>
	* includes DATAPRTP and all structure factor entries NO	NST1TP through NONS	ST10TP.
PDBPGMTP	<u>6250cpi</u> <u>1600cpi</u> <u>TK50</u>	<u>1/4"</u> <u>8mm</u>	<u>4mm</u>
(all computer programs and miscellaneous files)	VAX/VMS <i>copy</i>		0[]\$354
STRUCTURE FACTOR ENTRIES (experimental diffraction data)	[] NONST5TP[] NONST6TF [] NONST9TP[] NONST10T Choose one format for selection(s) shown above:	[]\$371[]\$34	<u>4mm</u> 0[]\$354
	* includes DATAPRTP and all structure factor entries NOI		1101P.
NMRRS1TP (NMR experimental data entries)	6250cpi 1600cpi <u>TK50</u> VAX/VMS backup[]\$359[]\$359[]\$421 VAX/VMS copy	<u>1/4" 8mm</u> []\$371[]\$340	<u>4mm</u> D[] \$354
	TOTAL CHARGES		
Foreign air mail charges (\$19 per tape item mailed outside U.S. and Canada) led from \$19.00 foreign air mail charges)	\$19.00	(if applicable)
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BROOKHAVEN ORDER FORM

PRINTED DOCUMENTATION (no charge) [] Atomic Coordinate and Bibliographic Entry Format Description for DATAPRTP (Feb. 1992) [] Complete List of Bibliographic Entries [] Current DATAPRTP Directory [] Data Deposition Form [] Detailed Contents and Format Description for Each Structure Factor Entry [] Latest Newsletter [] Sources of Visual Aids for Macromolecular Structure (Feb. 1990) [] User Guide (Summer 1992) [] Full Tables

PLACING AN ORDER (Prices are valid until September 30, 1993)

We *must* receive the following three items before service is provided (it is best to send all items together *via* mail -- facsimile orders are not acceptable):

- 1. completed order form
- 2. mailing label indicating exact shipping address
- 3. payment (use one of the methods listed below):
 - ⇒ Check payable to <u>Brookhaven National Laboratory</u> in U.S. dollars and drawn on a U.S. bank. Foreign checks are not acceptable.
 - ⇒ Original hardcopy of purchase order payable to <u>Brookhaven National Laboratory</u>. After your order is processed, our Fiscal Division will invoice you.
 - Wire transfer. In order to use wire transfer capabilities, we must <u>first</u> receive an <u>original purchase order from you</u>. After you receive our invoice, your bank should send a wire transfer to:

Bank name: Morgan Guarantee Trust Company of New York

Acct. name: Brookhaven National Laboratory

Cust. Acct.: 076-51-912

Please mail all required items to:

Protein Data Bank Orders Chemistry Department, Building 555 Brookhaven National Laboratory Upton, NY 11973 USA

Affiliated Centers

Eleven affiliated centers offer DATAPRTP for distribution. These centers are members of the Protein Data Bank Service Association (PDBSA). Centers designated with an asterisk(*) distribute DATAPRTP on magnetic media; those without an asterisk are online DATAPRTP distributors.

CAN/SND

Canadian Scientific Numeric Data Base Service Ottawa, Ontario, Canada Roger Gough 613-993-3294 cansnd@vm.nrc.ca

CAOS/CAMM

Dutch National Facility for Computer Assisted Chemistry Nijmegen, The Netherlands Jan Noordik 31-80-653386 noordik@caos.caos.kun.nl

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Prophet

BBN Systems and Technologies a division of Bolt Beranek and Newman Inc. Cambridge, Massachusetts Carl Foeller 617-873-2669 prophet-info@bbn.com

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Statement of Support

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