

Sequence Data Submission and Accession Numbers

Authors should be aware that all new nucleotide sequence data must be submitted together with a completed data submission form to the EMBL Nucleotide Sequence Database. Upon receipt, a unique database accession number to identify the sequence will be provided to the author within 2-3 working days. The database accession-number is included in the manuscript and allows the community to retrieve the data upon reading the journal article.

Nucleotide sequence data reported in FEBS Letters with an EMBL Nucleotide Sequence Database accession number can be retrieved electronically by sending commands to a File Server at the European Bioinformatics Institute. This service is free of charge. For example, to get the sequence with accession-number X55651, an electronic mail message should be sent to the Internet address NETSERV@EBI.AC.UK including the following line:

```
GET NUC:X55651
```

The requested data will be returned automatically via electronic mail.

The most important file server command, to get users started, is:

```
HELP
```

The file server offers various other services (e.g. documentation, submission form, access to nucleotide and protein sequence data, protein structure data, software etc), details of which are provided in the HELP file.

How to contact the EMBL Nucleotide Sequence Database

Postal address: EMBL Nucleotide Sequence Submissions
European Bioinformatics Institute
Hinxton Hall, Hinxton
Cambridge CB10 1RQ, UK.

Computer Network: DATASUBS@EBI.AC.UK (for data submissions)
ATALIB@EBI.AC.UK (for general enquiries)
UPDATE@EBI.AC.UK (for updates/publication notifications)
NETSERV@EBI.AC.UK (file server)

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