

BBA Short Sequence-Papers

In addition to Regular Papers, BBA's 'Gene Structure and Expression' section publishes Rapid Reports and, included in these, Short sequence-papers. General information pertaining to Rapid Reports can be found in the separate Announcement published regularly in BBA.

Short sequence-papers include data on structural RNA and their genes, DNA sequences coding for a protein, DNA sequences coding for regulatory regions (promoters, enhancers) and other DNA regions of general interest, like repetitive DNA. Intron sequences without known function, or comparative sequences of already well characterized genes are acceptable only if considered to be of sufficient interest.

Short sequence-papers should conform to the style of a Rapid Report, i.e., should not exceed four printed pages, including figures and tables. They should be organized as follows: Title (including names and affilia-

tions of authors, and key words); Summary; Introduction, results, discussion and acknowledgements in a continuous style, not exceeding 1000 words (about 1.5 printed pages); References.

A 'Short sequence-paper' should be submitted to Amsterdam. (Submission to any of the other centres will delay publication.) From the submission letter it should be clear that it is a 'Short sequence-paper' and the manuscripts should be marked 'Short sequence'.

Sequences must be presented in photographic (i.e., camera-ready) form, and should be deposited at the EMBL Data Bank (Heidelberg) **before submission**. The EMBL accession number should be mentioned in the submission letter and/or in a footnote to the paper. To accelerate publication, reviewing will be performed by a Managing Editor, thus ensuring a quick decision, generally within 2 weeks. Proofs will not be sent.