

1. 期刊基本信息

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2. 期刊简介 About the journal

Molecular Biology and Evolution



Molecular Biology and Evolution (MBE) publishes research at the interface between molecular and evolutionary biology. The journal publishes investigations of molecular evolutionary patterns and processes, tests of evolutionary hypotheses that use molecular data, and studies that use molecular evolutionary information to address questions about biological function at all levels of organization. Reports of work on comparative and evolutionary genomics and the evolution of molecular structure and function are particularly welcome.

MBE is entirely owned by [The Society for Molecular Biology and Evolution](#). The [Council](#) of the Society appoints the [Editor](#). The Journal is published by [Oxford University Press](#) in conjunction with [HighWire Press](#).

3. 投稿指南 Instructions to Author

All material to be considered for publication in *Molecular Biology and Evolution* should be submitted in electronic form via the journal's online submission system (<http://mbe.manuscriptcentral.com>). Full instructions for online manuscript submission can be found [here](#) or via the [online submission site](#). If for exceptional reasons you are unable to submit your manuscript via the online submission system, please contact the [Editorial Office](#).

Editorial policy

MBE welcomes manuscripts reporting both empirical and theoretical work, as well as significant new statistical and computational methods. All work must have a solid biological basis. *MBE* is a primary research journal, but review articles are also published.

***MBE* will only publish work focussed on taxonomy and systematics or descriptive studies of genetic diversity and population structure, if they are of interest and relevance to a broad audience.**

The decision to accept or reject a manuscript will be made as rapidly as possible. *MBE* receives many more manuscripts than can be published. To be accepted for publication, a manuscript must make a substantial contribution to the field and be of interest to a broad audience. Manuscripts that report work not suitable for publication in the journal may be returned to authors without detailed review. Authors may seek advice from the [Editor](#) on the suitability of a manuscript before submission.

Note that publication in *MBE* incurs [page charges](#) that must be paid in full before an article is published. We do not, however, refuse publication to authors who cannot pay charges. Contact the [Editorial Office](#) if you think this applies to you.

Manuscripts are considered for publication in *MBE* on the understanding that authors have complied with all ethical and privacy guidelines and/or legislation covering the work being reported. Manuscripts will only be considered for publication in *MBE* that are not under consideration for publication elsewhere. Any manuscript or any part of a manuscript that has been published or submitted for publication elsewhere cannot be accepted.

Corresponding authors are required to sign an agreement licensing use of their material to the Society for Molecular Biology and Evolution before the paper can be published. Material published in the journal cannot be reproduced or published elsewhere (including on the Internet) without the written permission of the copyright owner.

Publication is taken to imply that the authors are prepared to make available, on request and at reasonable cost, any strains, cell lines, or clones used in reported experiments, and any public-domain computer programs on which the reported work is based. Authors may be required to provide computer programs as part of the review process. Substantial programs that have not previously been published must be provided for publication as Supplementary Material, unless they are already available through the World Wide Web, in which case a URL must be supplied. Newly reported nucleic acid and amino acid sequences, and structural coordinates must be submitted to appropriate public databases (e.g., GenBank <http://www.ncbi.nlm.nih.gov/>; the EMBL Nucleotide Sequence Database <http://www.ebi.ac.uk/embl/index.html>, DNA Database of Japan <http://www.ddbj.nig.ac.jp/>; the Protein Data Bank <http://www.rcsb.org/pdb/>; Swiss-Prot <http://www.ebi.ac.uk/swissprot/>).

Proofs and offprints

The corresponding author will receive a unique URL that gives access to the electronic version of their published paper free of charge. Order forms for purchasing print offprints will be sent to the corresponding author with page proofs. Requests for electronic access and print offprints by other authors should be channelled via the corresponding author.

Page charges

There is a charge of \$40 per page or fraction of a page for the first ten pages (~30% of production cost). For each page over ten, the charge will be \$80 per page.

Articles may be held back from publication until page charges are paid in full. To avoid publication delay, arrange payment as soon as you receive an invoice. Payment can be made by check or credit card, but Oxford University Press regrets that it is unable to accept purchase orders.

Authors can ask the Editor to waive charges at the time a manuscript is accepted for

publication if NONE of the authors has ANY funds (grant, research allocation, departmental, or otherwise) to pay them. Late requests will not be considered. In any case, page charges will only be waived for the first ten pages. Waiving page charges shifts the cost of publication to other authors. For this reason charges are seldom waived.

Journal page numbers can be estimated as: $\text{Journal pages} = (0.3 \times \text{manuscript text pages}) + (0.44 \times \text{figures}) - (0.15 \times \text{tables})$ (Text pages include: title page, abstract, main text, literature cited, all table pages, and figure legends.)

Color figures will incur a charge of \$750 per figure. Charges for color figures cannot be waived, nor will page charges be waived for manuscripts that include color figures. Color versions of figures that appear in black and white in the journal can be published as online supplements. These are permanently linked to the article in the online journal and do not incur any additional cost.

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MANUSCRIPT PREPARATION

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**IT IS IMPORTANT TO FOLLOW THESE INSTRUCTIONS CAREFULLY.
FAILURE TO DO SO MAY DELAY PUBLICATION OF YOUR
MANUSCRIPT.**

Manuscript submission is now exclusively [online](#).

Note: Short communications are no longer published as "Letters to the Editor." Short manuscripts should be formatted in the same way as full-length manuscripts. They must contain new and previously unpublished data, analysis or theory. Manuscripts that contain only opinion, commentary or discussion will not be accepted.

Manuscripts must:

- conform to the guidelines in the *Council of Biology Editors Style Manual* (6th ed., 1994) <http://www.councilscienceeditors.org/>, and to the style used in recent issues of this journal and to the terminology and style conventions described below
- be concise. Long articles attract more page charges; submit

large tables, sequence alignments etc as online supplements to keep page numbers down

- be written in American English
- be organized in the sequence described below
- be double-spaced (revised manuscript resubmission only)
- have 25-mm page margins
- be typed no smaller than 12-point Times Roman
- have correct diacritics for non-English words
- include accession numbers for all newly reported sequences and structural coordinates
- acknowledge all funding sources
- be saved as a .doc, LaTeX, TeX, .rtf, .ps or .pdf file
- include text, tables, and figure legends in a single file
- include figures as separate files.

Keep formatting simple:

- avoid hyphenation, justification, linked and embedded objects and images, and other advanced word processing features, as they have to be removed during typesetting
- avoid unusual fonts: stick with Symbol, Times/Times New Roman, Courier/Courier new, Helvetica/Arial
- be consistent in representing symbols
- distinguish similar-looking symbols (e.g., letter x, multiplication sign, and Greek chi; minus sign, hyphen, and dash)
- do not approximate characters by creating your own symbols (e.g., superscript o for degree symbol)
- do not approximate formatting (e.g., linebreak + tab for hanging indent)
- do not use underlining to indicate italics or in plus-minus signs

Title Page

must contain:

- title
- names of all authors
- institution(s) at which research was done
- current affiliations of all authors
- name and address, telephone number, fax number, and e-mail address of the corresponding author
- list of 3 to 6 key words
- running head (maximum of 50 characters and spaces)
- list of nonstandard abbreviations used, if any (see terminology and style conventions below)

Choose a title that contains useful information about content rather than one with dramatic impact. It will be more accessible to readers searching PubMed and other electronic databases.

Choose key words that accurately index your article to a broad readership. Include essential words that also appear in the title, as it may be searched separately. Species names should be among the keywords in articles based on a single species.

Abstract

The abstract should be a factual condensation of the entire paper, including a statement of purpose, a clear description of observations and findings, and a concise presentation of conclusions.

The abstract must not exceed 350 words (one double-spaced page). References to cited literature should not be included.

The main text should comprise:

- Introduction
- Materials and Methods
- Results
- Discussion (can be combined with Results)
- Conclusion (if needed)
- Supplementary Material (i.e., indication of availability where appropriate)
- Acknowledgments
- Literature Cited

Literature cited within the text must:

- be cited by author and year
- include relevant pages for quotations
- be in chronological order when grouped (alphabetical order if published in the same year)
- include the names of all authors when there are three or fewer authors
- include the name of the first author and "et al." when there are four or more authors
- refer to unpublished work of authors as "unpublished data"
- refer to unpublished work of others as "name, personal communication" (permission to cite is required)

The 'Literature Cited' section must

- be arranged alphabetically by author(s) and then chronologically
- contain only works specifically cited in the text
- refer to manuscripts accepted for publication but which have not yet been published as "in press"
- not include manuscripts that have not formally been accepted for publication
- follow the guidelines for abbreviations of periodicals in *The Council of Biology Editors Style Manual* (6th ed., 1994) <http://www.councilscienceeditors.org/index.php3>
- be formatted as in recent issues of the journal and as follows:

Journal articles:

Pensole, G., C. Gissi, C. Lanave, and C. Saccone. 1995. Glutamine synthetase gene evolution in bacteria. *Mol. Biol. Evol.* **12**:189-197.

(> 10 co-authors):

Wilson, R., R. Ainscough, K. Anderson et al. (53 co-authors). 1994. 2.2 Mb of contiguous nucleotide sequence from chromosome III of *C. elegans*. *Nature* **368**:32-38.

Books:

Ingram, V. M. 1963. *The hemoglobins in genetics and evolution*. Columbia University Press, New York.

Book chapters:

Hall, B. G. 1983. Evolution of new metabolic functions in laboratory organisms. Pp. 234-257 *in* M. Nei and R. K. Koehn, eds. *Evolution of genes and proteins*. Sinauer Associates, Sunderland, Mass.

Supplementary material

When appropriate, material such as sequence alignments and large tables can be published online as supplementary material permanently linked to an article in the online journal. The **Supplementary Material** section should only list any supplementary material available online at the MBE [Web site](#). The material itself should be prepared as described below in "Preparation of supplementary material."

Tables

Tables must:

- have short titles that describe their contents
- not have "legends"
- not include vertical rules
- continue onto a second or third page, if necessary.
- be set in a font equivalent in size to 12 point Times Roman type.
- be numbered with Arabic numerals
- arrange data so that columns of like material read down, not across
- include sufficient information that the meaning of the data is clear without reference to the text
- use abbreviations to conserve space
- include explanatory material as footnotes immediately below the table
- reference explanatory footnotes by superscripted lower case letters except for significance levels, for which asterisks are used
- not include detailed descriptions of experiments in explanatory footnotes
- be used only when six or more individual data are presented

Figure legends

Figure legends must:

contain enough information for the figure to be understood independently of the text

- define abbreviations used in the figure

All items from 'Title page' to here **must** be saved in a single file.

Figures

For useful information on preparing your figures for publication, go to <http://cpc.cadmus.com/da>.

Figures must:

- be numbered consecutively following the sequence in which they are mentioned in the text
- include scale bars where appropriate. These should not be placed in the legend
- be the size intended for publication. Maximum single column width = 8.2 cm/3.25 inches; 2 columns = 16.9 cm/6.75 inches; depth = 24 cm/9.25 inches
- have uniform lettering style
- not include sequences and alignments that exceed one journal page. Larger alignments must be submitted as online Supplementary Material at the **MBE** Web site and their availability indicated in print in the **Supplementary Materials** section

Programs suitable for creating production-ready graphics files are the latest versions of:

- Adobe Photoshop (latest version)*
- Adobe Illustrator (latest version)*
- Canvas (version 7 or later)
- Corel Draw (version 7 or later)
- Macromedia FreeHand (version 8 or later)

* Preferred

Electronic versions of figures must:

- use Adobe Postscript fonts, NOT TrueType or system "bitmap" fonts
- be converted to grayscale or bitmap mode if originally in color but to be reproduced in black and white
- have any unnecessary white space cropped from around the outside of the image if the file is a TIFF file. Be careful not to crop any of the intended image
- each be saved in a separate file. If a figure has multiple parts (e.g., Fig. 1A, Fig. 1B) all parts should be saved into one file
- be saved at 1200 dpi for line figures and 350 dpi for half tones
- for publication, be saved as .tif or .eps files
- always use the latest version of the software program available. Files from older versions often lose integrity when opened in newer versions.

Files created using Microsoft PowerPoint are not suitable for production. If you are not able to provide figures in one of the above formats please contact the [Editorial Office](#).

Color figures must:

- be in CMYK mode NOT RGB mode
- be saved at 350 dpi
- **be paid for.** Color figures will incur a charge of \$750 per figure. Charges for color figures cannot be waived, nor will page charges be waived for manuscripts that include color figures. Figures can be published in black and white (normal page charges apply) in the article, with a color version published as supplementary material permanently linked to the online journal. There is no cost for publishing online supplementary material.

Preparation of supplementary material

All material to be considered as supplementary data must be submitted at the same time as the main manuscript for peer review. Please indicate clearly the material intended as Supplementary Material upon submission. Also ensure that the supplementary material is referred to in the main manuscript at an appropriate point in the text. It must be supplied to the production department with the article for publication, not at a later date. It cannot be altered or replaced after the paper has been accepted for publication.

Files for supplementary material should be clearly marked as such, and be accompanied by a summary of the file names and types.

Please note that supplementary material will not be copyedited, so ensure that it is clearly and succinctly presented, and that the style of terms conforms with the rest of the paper. Also ensure that the presentation will work on any Internet browser.

A maximum of 5 files is acceptable to make up the supplementary material unit for an article. The maximum size per file should not exceed 2 MB (though text files should be a great deal smaller), and files must be as small as possible, so that they can be downloaded quickly. An HTML index page is usually created to link the supplementary data file(s) to the article. Please provide short (24 word) titles for each individual file; these will be used to create links to the files from the index page.

The following formats are acceptable:

- Plain text (.txt)
- HTML (.html, htm)
- Jpeg (.jpg, .jpeg)
- GIF (.gif)
- QuickTime video (.mov)
- MPEG Movie (.mpg)
- Microsoft AVI Video (.avi)
- Adobe PDF (.pdf)
- Microsoft Excel Spreadsheet (.xls)

Sequence alignments should be submitted as supplementary material in two forms:

1. As a text file in a "sequential" format such as fasta, so that readers can download and use the alignment.
2. In an "aligned" or "interleaved" format, so that readers can easily view the alignment.

Links from articles to online supplementary material will be inserted as soon as articles are published online (approximately one week ahead of the print journal).

Sequence alignments can also be deposited at EMBL-ALIGN:
http://www.ebi.ac.uk/embl/Submission/align_top.html.

Submission of accepted manuscripts

Authors will be notified of receipt of their manuscript by the online submission system. The Editor, [William Martin](#), is responsible for the peer review process and for deciding whether the manuscript should be accepted, returned for revision, or rejected. The corresponding author will be able to view the stage of processing of the manuscript at the submission system Web site. Authors will receive a final decision e-mail message from the Editor. If, at any time, you are dissatisfied with any aspect of the handling of your manuscript, or if you have any suggestions that might improve our performance, please contact the Editor.

Following review, resubmit revised manuscripts [online](#). **These must be submitted *both*:**

- **in a single .pdf file including all the components of the paper (The .pdf file is essential to ensure that illustrations can be checked for corruption in transit and to enable your paper to be processed for online MBE Advance Access. Supplementary material should be submitted as separate pdf file(s) for checking but will not normally be included in MBE Advance Access.) *and***
- **as .doc, .rtf, TeX, or LaTeX (for text), .tif or .eps files (for illustrations), and as listed above under Preparation of Supplementary Material (for supplementary materials). These are the files that will be used for production of the final printed and electronic files of your paper. Note that for production purposes tables must be double-spaced, each beginning on a new page, and that figure legends should be grouped together starting on a new page and should also be double spaced.**

(JPEG and GIF files, and files created using Microsoft PowerPoint, are not suitable for production.) Contact the [Editorial Office](#) if you are not able to provide figures in one of the acceptable formats.

All files must be given names that include BOTH the manuscript handling number AND the corresponding author's name. The

manuscript submission system automatically allocates manuscript numbers of the form MBE-03-0058, where 03 is the year of submission and 0058 the manuscript number. Files pertaining to ms MBE-03-0058 submitted by author Chen should therefore be named Chen03-0058.pdf; Chen03-0058.rtf; Chen03-0058.eps; Chen03-0058.html, etc.

Terminology and style conventions

- **Abbreviations and symbols** should follow the International System of Units (SI)
<http://physics.nist.gov/Pubs/SP330/contents.html>.
- **Nonstandard abbreviations** must be defined at first occurrence, in both abstract and main text.
- **Species** must be identified by italicized scientific binomens. **Generic names** that are also common names should not be italicized unless they form part of a binomen (thus an investigation may involve *Drosophila melanogaster* or *D. melanogaster*, but a comparison would be made between *Drosophila* and human genes). Binomen abbreviations of the form "Gsp" (Genus, species) as, for example, "Hsa" for *Homo sapiens* may be used in tables and figures.
- **Genetic loci** must be italicized and must follow the established rules of genetic nomenclature that have been established for the various organisms (e.g. HUGO Nomenclature Committee
<http://www.gene.ucl.ac.uk/nomenclature>, the International Immunogenetics Database <http://imgt.cnusc.fr:8104/>, Mouse Nomenclature Guidelines & Locus Symbol Registry <http://www.informatics.jax.org/support/nomen/>, Mendel-GFDb (plants) http://jii06.jic.bbsrc.ac.uk/gfd_index.html
- The International Union of Pure and Applied Chemistry and the International Union of Biochemistry and Molecular Biology rules of nomenclature (<http://www.chem.qmw.ac.uk/iupac/bibliog/white.html>) must be followed for **amino acids, peptides, proteins, enzymes, nucleic acids, polynucleotides, carbohydrates, and lipids.**
- **Mathematical equations** must be presented with correct spacing between characters. Characters in equations and their counterparts in the text will be set in italics unless the author specifies otherwise the first time a character appears. Equations must be numbered sequentially, in Arabic numerals in parentheses, on the right-hand side of the page.
- **Statistical inferences**, including those used for phylogenetic estimation, must be soundly based. Where appropriate, the assumptions underlying statistical inferences should be made explicit and sources of error should be clearly identified.
- **The IUBMB single-letter code for nucleotide bases** including ambiguity is as follows: A = adenine; C = cytosine; G = guanine; T = thymine; U = uracil; R = A/G (purine); Y = C/T (pyrimidine); M = A/C; W = A/T; S = C/G; K = G/T; B = C/G/T (not A); D = A/G/T (not C); H = A/C/T (not G); V = A/C/G (not T); and N = X = A/C/G/T (any or unknown). For ambiguous nucleotides, T and U are equivalent.

Effective interdisciplinary communication requires that the precise meaning of words be understood in both disciplines. The following conventions should generally be followed.

1. When aligned sequences are compared, differences, not changes, may be observed. Changes can only be inferred, and a single difference may result from multiple changes.
2. The word "invariant" has two common but different meanings, invariable and unvaried. Either can be used, but the meaning must be clear.
3. A mutation generally occurs in a single individual and gives rise to an allele. If an allele achieves some frequency in a population it can be referred to as a polymorphism (not a "common [or rare] mutation"). If it has become fixed in a population it may be referred to as a substitution.
4. If two molecules are alike in some degree, they are similar. If it is inferred from their similarity that they have a common ancestor, then they are homologous, but if their similarity was acquired by convergence, they are analogous. When homology arises via a gene duplication (all or part), it is paralogy; when it arises via speciation, it is orthology; when it arises by horizontal gene transfer, it is xenology.
5. The phrase "insertions and/or deletions" may be reduced to "indels."
6. Gaps are introduced into sequences to increase their similarity rather than to optimize similarity, unless an algorithm is employed that guarantees an optimized result according to the way similarity is defined (e.g., as maximum matches).
7. Similarity should not be asserted to be significant unless accompanied by a probability statement and its method of determination.

Nucleotide sequences

- Newly reported sequences must be deposited in the DDBJ/EMBL/GenBank database. Accession numbers must be included in the final version of the manuscript and cannot be added at the proof stage.
- Nucleotide sequences must be accurate. Measures employed to avoid error should be clearly stated. Determining the sequence on both strands is the standard for most studies. When only one strand is sequenced, this must be explicitly stated both in Materials and Methods and in the entry that is submitted to one of the sequence databases.
- All ambiguous positions in a nucleotide sequence must be indicated with the appropriate IUBMB single-letter code (see "terminology" above), rather than "resolved" by guesswork.

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INSTRUCTIONS FOR MANUSCRIPT SUBMISSION

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Follow the [manuscript preparation](#) instructions carefully. Failure to follow these instructions may lead to delay in the publication of your manuscript. Use version 4.0 or higher of Netscape or Internet Explorer. If you have trouble finding manuscripts or have other problems with your account do not create another account. Instead, e-mail support@scholarone.com or telephone +1 434 817 2040 extension 167.

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 - Select Browse to locate your files. Select the file type from the pull-down menu. An initial submission of the main manuscript should be uploaded as a single .pdf file containing all components of the manuscript including text, tables and figures. (Supplementary Material files should be uploaded as separate files formatted as they will appear if accepted for publication.) Initial submission as a single .pdf file greatly facilitates the review process and is strongly encouraged. If you are asked to revise and resubmit your manuscript, you will be REQUIRED to prepare your manuscript both as a single .pdf file and as files suitable for journal production (see [Manuscript preparation](#)), before it can be accepted for publication. If for any reason you are unable to prepare your manuscript as a single .pdf file in the first instance, submit the text as a .doc, .rtf or .ps file, which will be converted to .pdf format, and submit images as separate .gif, .jpg, .eps, .png, and .tif files. These will be converted to small .jpg files. Converted .pdf and .jpg files will be the files evaluated during the review process.
 - Select whether files should be considered for review, (default is Yes). If you select No, editorial staff will still be able to view the file and make it available to an editor or reviewer if necessary.

- Select Upload to submit your file.
 - When an upload is complete, you will see a confirmation window asking for a description of the file. Refer to your Main Document as Manuscript Text or Main Document. Refer to figures by number, e.g. Figure 1 or Fig. 1. For other supporting files indicate clearly what the file is as well as its format (MS Excel, MS Word, etc.).
3. The on-screen version of your manuscript, which is the version accessed by the editor and reviewers, can be viewed and proofed after it has been successfully uploaded.
 4. You can remove files and repeat the upload process at the File Upload screen, accessed by selecting Previous.
 5. When you are satisfied with the uploaded manuscript select Submit. It is not until this button is pushed that the manuscript and all of the associated information (i.e., contributing authors, institutions, etc.) is linked together and the manuscript is given a manuscript number. Once the manuscript is submitted it is not possible to undo the submission.
 6. After the manuscript has been submitted you will receive an e-mail confirmation stating that your manuscript was successfully submitted. This e-mail will also give the assigned manuscript number, which is used in all correspondence. If you do not receive this e-mail, your manuscript will not have been successfully submitted to the journal and the paper cannot progress to peer review. If this is the case your manuscript will still be sitting in the Partially Submitted Manuscripts section of your Author Centre awaiting your attention.
 7. If you return to your Author Centre you will notice that your newly submitted manuscript can be found in the Submitted Manuscripts area. Among the information listed there, the Processing Status section provides information on the status of your manuscript as it moves through the review process.

SUBMITTING A REVISED MANUSCRIPT

1. Log on to the [Manuscript submission](#) site as before. Access your manuscript by selecting its title in the Manuscripts to be Revised section of the Author Centre.
2. Upload your revised manuscript using the File Manager screen. It is essential that you upload your revised manuscript as a .doc .rtf or .ps file. PDF files cannot be used for production.
3. Paste your response to reviewers' and editor's comments in the text areas at the bottom of the View comments/respond screen.

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4. 编辑部信息 Editorial Board

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