

Special Issue

Ancient DNA and Molecular Archaeology

Message from the Guest Editors

Ancient DNA (aDNA) is any DNA extracted from ancient specimens, important for diverse evolutionary research. During the last three decades, DNA analysis on degraded samples has revealed itself as an important research tool in anthropology, molecular evolution, and population genetics. The past decade has witnessed a revolution in ancient DNA research. Although the field's focus was previously limited to mitochondrial DNA and a few nuclear markers, whole-genome sequences from the deep past can now be retrieved. This breakthrough is tightly connected to the massive sequence throughput of next-generation sequencing platforms and the ability to target short and degraded DNA molecules, with the possibility to go back in time 400,000 years for samples from temperate regions and 700,000 years for permafrozen remains. This Special Issue welcomes original research, brief research reports, and review papers about molecular archaeological studies on unearthed materials (including animals, plants, humans, sediments, and so on), using genome-wide sequencing or microarray technologies or various kinds of markers, such as mtDNA, Y chromosome, STRs, SNPs, and isotopic data.

Guest Editors

Dr. Shaoqing Wen

Institute of Archaeological Science, Fudan University, Shanghai 200433, China

Dr. Pengfei Sheng

Institute of Archaeological Science, Fudan University, Shanghai 200433, China

Deadline for manuscript submissions

closed (20 March 2025)

G C A T
T A C G
G C A T

Genes

an Open Access Journal
by MDPI

Impact Factor 2.8
CiteScore 5.5
Indexed in PubMed



mdpi.com/si/214799

Genes
Editorial Office
MDPI, Grosspeteranlage 5
4052 Basel, Switzerland
Tel: +41 61 683 77 34
genes@mdpi.com

[mdpi.com/journal/
genes](https://mdpi.com/journal/genes)



G C A T
T A C G
G C A T

Genes

an Open Access Journal
by MDPI

Impact Factor 2.8
CiteScore 5.5
Indexed in PubMed



[mdpi.com/journal/
genes](https://mdpi.com/journal/genes)



About the Journal

Message from the Editor-in-Chief

Genes is central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fast-moving field. There is a need for good quality, open access journals in this area, and the *Genes* team aims to provide expert manuscript handling, serious peer review, and rapid publication across the whole discipline of genetics. Starting in 2010, the journal is now well established and recognised. Why not consider *Genes* for your next genetics paper?

Editor-in-Chief

Prof. Dr. Selvarangan Ponnazhagan
Department of Pathology, The University of Alabama at Birmingham,
1825 University Blvd, SHEL 814, Birmingham, AL 35294-2182, USA

Author Benefits

Open Access:

free for readers, with article processing charges (APC) paid by authors or their institutions.

High Visibility:

indexed within Scopus, SCIE (Web of Science), PubMed, MEDLINE, PMC, Embase, PubAg, and other databases.

Journal Rank:

JCR - Q2 (Genetics and Heredity) / CiteScore - Q2 (Genetics (clinical))