

Special Issue

Feature Papers in the Population and Evolutionary Genetics and Genomics Field 2026

Message from the Guest Editors

This Special Issue seeks to showcase outstanding contributions that advance conceptual, methodological, and empirical research in the field. We aim to highlight studies that elucidate the genetic and genomic mechanisms underlying population structure, evolutionary change, and speciation.

We particularly encourage submissions that integrate high-throughput sequencing technologies, genome-wide analyses, quantitative modeling, and interdisciplinary perspectives to address fundamental and emerging questions in evolutionary biology. Contributions bridging theoretical frameworks with empirical data, or linking genomic variation to ecological, phenotypic, and environmental contexts, are especially welcome.

We invite original research articles and comprehensive reviews covering, but not limited to, population structure and demographic inference, patterns and determinants of genetic diversity, evolutionary genomics and genome-wide variation, natural and sexual selection across spatial and temporal scales, local adaptation and adaptive evolution, speciation, hybridization, and reproductive isolation, quantitative, functional, and comparative genomics in evolutionary contexts, etc.

Guest Editors

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Deadline for manuscript submissions

31 December 2026

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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?

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