

Special Issue

Cotton: Genomics, Biotechnology and Molecular Breeding

Message from the Guest Editor

Benefited from the fast progress of cotton genomics, many QTL/genes have been identified by linkage mapping and genome-wide association analysis, and functional research of genes has also sped up. With the re-sequencing of thousands of germplasms, many markers associated with important traits have been discovered, which will facilitate the molecular breeding of cotton. Also, new biotechnologies, such as CRISPR and genotype-independent cotton genetic transformation, have been developed. In order to exhibit the latest progress in cotton, we welcome cotton scientists to submit their new findings in *Biology*.

Guest Editor

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Deadline for manuscript submissions

closed (31 May 2025)



Biology

an Open Access Journal
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Impact Factor 3.5
CiteScore 7.4
Indexed in PubMed



mdpi.com/si/207480

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Message from the Editorial Board

A major strength of biological science is the diversity of approaches that biological scientists apply to their research problems. *Biology* reflects this diversity and brings together studies employing the varied experimental and theoretical approaches that are fueling biological discovery. *Biology*, the journal, is a fully peer-reviewed publication with a rapid and economical route to open access publication and is listed on PubMed. All articles are peer-reviewed and the editorial focus is on determining that the work is scientifically sound rather than trying to predict its future impact.

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