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DESCRIPTION

Protist is the international forum for reporting substantial and novel findings in any area of research on protists. The criteria for acceptance of manuscripts are scientific excellence, significance, and interest for a broad readership. Suitable subject areas include: molecular, cell and developmental biology, biochemistry, systematics and phylogeny, and ecology of protists. Both autotrophic and heterotrophic protists as well as parasites are covered. The journal publishes original papers, short historical perspectives and includes a news and views section.

AUDIENCE

Researchers with an interest in protists working in the areas of protistology, protozoology, phycology, parasitology, molecular, cell and developmental biology, biochemistry, biotechnology, systematics, evolution and phylogeny, and ecology.

IMPACT FACTOR

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GUIDE FOR AUTHORS

INTRODUCTION

Protist publishes papers that report substantial and novel findings in any area of research on protists. Suitable subject areas include: molecular, cell and developmental biology, biochemistry, systematics and phylogeny, and ecology of autotrophic and heterotrophic protists. The journal will not, in general, publish floristic/faunistic accounts, descriptions or redescriptions of taxa which are not based on new original data, taxonomic work which lacks reproducibility or papers that are preliminary or merely confirmatory. The criteria for acceptance of manuscripts are scientific excellence, significance, and interest for a broad readership.

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Protist Genome Reports

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PGR will consider articles that announce a draft nuclear genome of a protist. Organelle (plastid, mitochondrion) genome sequence(s) will only be considered in combination with the draft nuclear genome sequence. The protist from which the nuclear genome sequence has been derived must be deposited in a public culture collection and the strain number provided before publication. The manuscript should provide a brief rationale for sequencing the particular protist including relevant references, with an emphasis on the detailed methodologies and protocols used in the assembly, annotation, and quality evaluation of the genome sequence. Accession numbers for both raw reads and assemblies, should be provided with the submission. The report will not be considered for publication until public availability of the sequence has been verified.

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Acknowledgements

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State the objectives of the work and provide an adequate background, avoiding a detailed literature survey or a summary of the results.

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Cohen J, Beisson J (1988) The Cytoskeleton. In Görtz HD (ed) *Paramecium*. Springer-Verlag, Berlin, pp 363–392

Margulis L, Corliss JO, Melkonian M, Chapman DJ, eds (1990) *Handbook of Protozoa*. Jones and Bartlett Publ, Boston, 914 p

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