***Genes & Genetic Systems***

**Instructions to Authors**

***Genes & Genetic Systems (GGS)*** is the official journal of the Genetics Society of Japan, which has been published bimonthly since 1918. ***GGS***, one of the most time-honored scientific journals in the world, is now an international peer-reviewed Open Access journal.

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# AIMS & SCOPE

*Genes & Genetic Systems*is an international peer-reviewed Open Access journal. It publishes important original papers and review articles bimonthly that address a wide range of fields in genetics. These include general genetics, molecular genetics, cell genetics, developmental genetics, behavioral genetics, ecological genetics, chemical genetics, epigenetics, cytogenetics, population genetics, evolutionary genetics, and molecular evolution. *GGS* also welcomes papers in the following fields, if they are related to genetics: genome-wide research, bioinformatics, systems biology, database, genetic bioresources, and technical advances (in both experimental and in silico methods). *GGS* maintains the highest standards of scientific and publication ethics. Articles are available freely online to a diverse global audience.

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Reviews and Minireviews summarize literature on a specialized topic, discuss any challenges in that field of research, and present a perspective on the future of the field. There is no minimum or maximum length for Reviews but Minireviews must be less than 3000 words (excluding References). Reviews and Minireviews are usually invited and managed by the Reviews Editor; recipients of the Genetics Society of Japan’s Young Scientist Award are also invited to collaborate with distinguished researchers to prepare a Minireview for consideration.

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## Acknowledgments

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*Journal articles*

Tajima, F. (1989) Statistical method for testing the neutral mutation hypothesis by DNA polymorphism. Genetics **123**, 585-95.

Saitou, N., and Nei, M. (1987) The neighbor-joining method: a new method for reconstructing phylogenetic trees. Mol. Biol. Evol. **4**, 406-425.

Sarojam, R., Sappl, P. G., Goldshmidt, A., Efroni, I., Floyd, S. K., Eshed, Y., and Bowman, J. L. (2010) Differentiating *Arabidopsis* shoots from leaves by combined YABBY activities. Plant Cell **22**, 2113-2130.

Sasaki, T., Matsumoto, T., Yamamoto, K., Sakata, K., Baba, T., Katayose, Y., Wu, J., Niimura, Y., Cheng, Z., Nagamura, Y., et al. (2002) The genome sequence and structure of rice chromosome 1. Nature **420**, 312-316.

*Books*

Steeves, T. A., and Sussex, I. M. (1989) Patterns in Plant Development. Cambridge University Press, Cambridge, UK.

Nakazaki, T., Naito, K., Okumoto, Y., and Tanisaka, T. (2008) Active transoposons in rice. *In* Rice Biology in the Genomics Era. (eds.: Hirano, H.-Y., Hirai, A., Sano, Y., and Sasaki, T.), pp. 69-79. Springer, Heidelberg.

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