

Feedback form

The subject of this module is continuously changing, making periodical updating desirable. For this reason we offer the module in electronic format. To help us with the task of updating, we would appreciate receiving relevant comments and requests from users to further improve the module. Your comments on this tool's usefulness will also be very helpful.

Below we have asked a few guiding questions for you to answer. You may send your answers to us directly by clicking the 'Send' button at the end of this page, by fax (no.: 57-2 445 0096), by post to your contact in IPGRI or by e-mail to either cdevicente@cgiar.org; cflb@lamolina.edu.pe; or tf12@cornell.edu

Questions:

1. Tell us how you use this tool...

2. Do you think the module:
 - a. Provides an overview of the basic scientific concepts of population genetic for assessing genetic diversity?

Yes
No
Why?

 - b. Shows how the use of molecular marker technologies can be a practical approach for anyone planning research on genetic diversity analysis.

Yes
No
Why?

 - c. Gives information on how to analyze and interpret molecular marker data.

 - d. Makes available a list of current bibliographical resources for each of the items discussed?

Yes
No
Why?

3. Tell us what you liked and what you would change about the following:
 - a. Contents, in general
 - b. Structure within the submodules
 - c. Drawings and diagrams
 - d. Appendices
 - e. Other

4. What would you add to the module to make it more useful in your particular situation?
 - a. Basic references
 - b. Examples of applications
 - c. Additional information on:
 - I. Concepts of population genetics
 - II. Measures of genetic diversity
 - III. Software described
 - d. Other

5. What other topics would you like to see in this module?

6. Was the module, or parts of it, difficult to download? Why?

Send