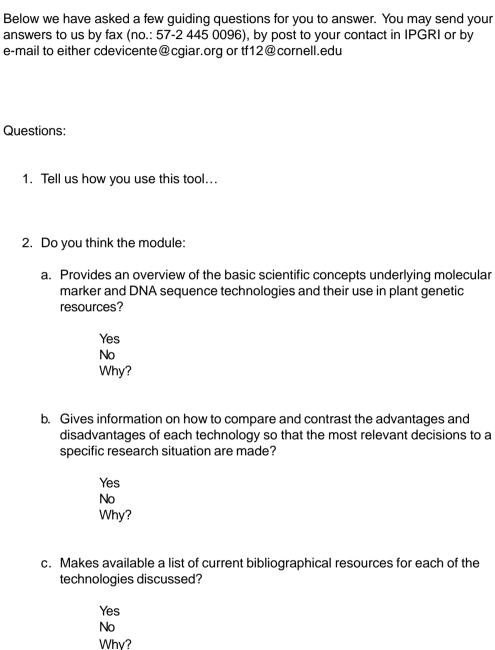# Feedback form

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 by fax (no.: 57-2 445 0096), by post to your contact in IPGRI or by e-mail to either cdevicente@cgiar.org 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 underlying molecular marker and DNA sequence technologies and their use in plant genetic resources?

        Yes
        No
        Why?

    b. Gives information on how to compare and contrast the advantages and disadvantages of each technology so that the most relevant decisions to a specific research situation are made?

        Yes
        No
        Why?

    c. Makes available a list of current bibliographical resources for each of the technologies discussed?

        Yes
        No
        Why?

3. Tell us what you liked and what you would change about the following:

   a. Contents

   b. Spectrum of technologies offered

   c. Criteria to select techniques

   d. Structure within the submodules

   e. Drawings, diagrams, pictures

   f. 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 basic concepts

   d. More technologies:

      i. Old

      ii. New

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

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