

Data submission

We are working towards on-line submission of your data through a web interface and automated mapping of your data against the dataset from the IBM high-resolution genetic map produced as a collaboration through NSF Grant # 9872655. To expedite data handling, be sure you use the data collection template included in the kit or downloadable from http://www.cafnr.missouri.edu/mmp/dna_kits.htm. In the interim please attach your map data in the data collection template and email it to IBMKits@missouri.edu. Be sure to include the control primer data you collected. This is required for us to process your data. It can be submitted as a separate file or within the same file as the novel map data. Your data will be mapped by a staff member.

Please attach the following additional information in a tab delimited text file:

Proposed Locus Name*	Primer sequence	What did type of sequence did you derive the primer from? cDNA, BAC end, etc.	GenBank number derived from if available	BAC number and library if appropriate

*These names will be checked against the current list in the Maize Genome Database to avoid unintentional duplications. In the event that a name already exists, we will contact you by email.

Include the following information in the body of your email message:

Your name:

Faculty whose lab is submitting the data:

Your postal address:

When this data can be made public: now, following publication, or in 1 year. If it's something other than these please explain.

Until the automated web interface is available mapping will be done in batches every two weeks. You will receive map position information by email within 3 weeks.

If you have questions about data submission, email IBMKits@missouri.edu. Due to the volume of materials,

phone inquiries of project personnel will not be accepted.