

## Pepper Genome Data Release Policy

**PLEASE READ BEFORE DOWNLOADING THE PRE-PUBLICATION PEPPER GENOME DATA:** We are pleased to make available the pre-publication pepper genome sequences of *C. annuum* var. *annuum* ‘Zhangshugang’, *C. baccatum* var. *pendulum* ‘PI 632928’ and *C. pubescens* ‘Grif 1614’, the *Capsicum* pan-genome graphs and the variant data of 500 *Capsicum* accessions for use by public and private research communities as a resource to facilitate plant biology discoveries and plant breeding programs.

The pre-publication pepper genome sequences, pan-genome graphs and variant data are made available with the understanding that users will respect the rights of those who contributed to this effort to describe the genomes, pan-genome and variant data in a peer-reviewed publication. By accessing these data, users agree not to publish any articles containing analyses of genes, genomic or variant data on a whole genome or chromosome scale prior to publication of the data by us. The restricted analyses include the identification of complete sets of genomic features such as genes, gene families, regulatory elements, repeat structures, etc., whole-genome comparisons of regions of evolutionary conservation or change, and population genomic analyses. Studies of any type on the reserved data sets that are not in direct competition with those planned by us may be undertaken following an agreement. Interested parties are encouraged to contact Dr. Zhangjun Fei (zjf25@cornell.edu) to discuss such possibilities.

The data may be freely used by all who respect the restrictions described above. Once moved to publication status, the genome sequences, pan-genome graphs and variant data are freely available for any subsequent use.