

# Gramene: A genomics and genetics resource for rice and other grasses.

C. Hubbard<sup>1</sup>, S. Avraham<sup>2</sup>, E. S. Buckler<sup>3,4</sup>, P. Canaran<sup>2</sup>, T. Casstevens<sup>3</sup>, B. Faga<sup>2</sup>, B. Hurwitz<sup>2</sup>, P. Jaiswal<sup>1</sup>, C. Liang<sup>2</sup>, S. McCouch<sup>1</sup>, J. Ni<sup>1</sup>, K. Ratnapu<sup>2</sup>, L. Ren<sup>2</sup>, S. Schmidt<sup>2</sup>, W. Spooner<sup>2</sup>, L. Stein<sup>2</sup>, I. Y. Teclie<sup>1</sup>, D. Ware<sup>2,4</sup>, S. Wei<sup>2</sup>, I. Yap<sup>1</sup>, K. Youens-Clark<sup>2</sup>, W. Zhao<sup>2</sup>  
(1) Department of Plant Breeding and Genetics, 240 Emerson Hall, Cornell University, Ithaca, NY, 14853, USA; (2) Cold Spring Harbor Laboratory, 1 Bungtown Rd, Cold Spring Harbor, NY, 11724, USA; (3) Institute for Genomic Diversity, Diversity, Cornell University, Ithaca, NY, 14853, USA; (4) USDA-ARS NAA Plant, Soil & Nutrition Laboratory Research Unit, Cornell University, Ithaca, NY, 14853, USA



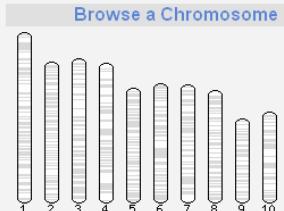
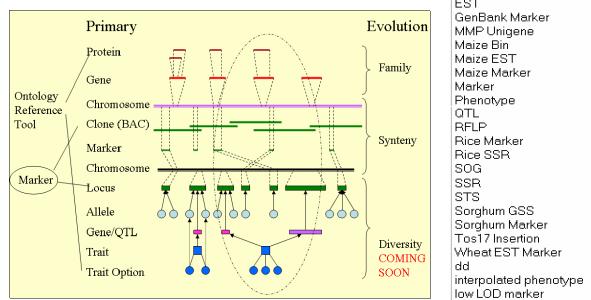
Breeders are concerned with the selection of favorable phenotypes. They have always endeavored to 'read' genome sequence by peering through the lens of the phenotype.

Measuring such characteristics by conventional methods is time-consuming and expensive, since it requires the organism to grow to maturity.

Hank Beachell at the Beaumont Experiment Station, TX, circa ~1935.

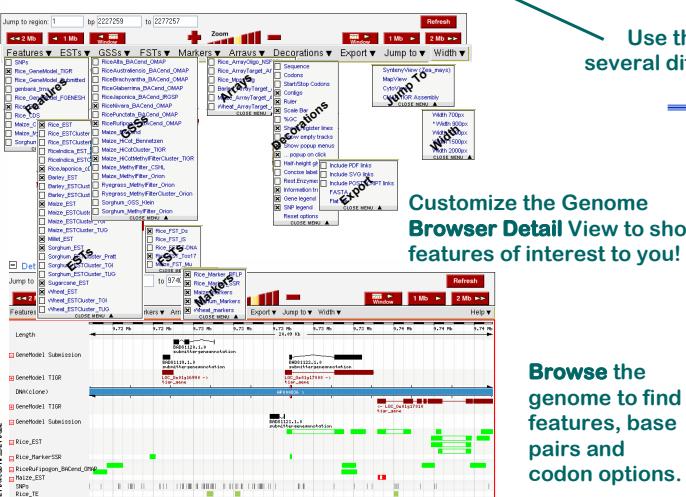
Molecular markers are the genetic signposts (DNA segments) that flag the presence of genes that control particular traits. Molecular markers offer a lens for identifying genes associated with phenotypes of interest.

Everything that can be mapped to the genome is called a "Feature."



In the Genome Browser, the rice genome's 12 chromosomes are displayed in a mapset of 12 maps.

Use the CMap viewer to compare 2 or more maps.

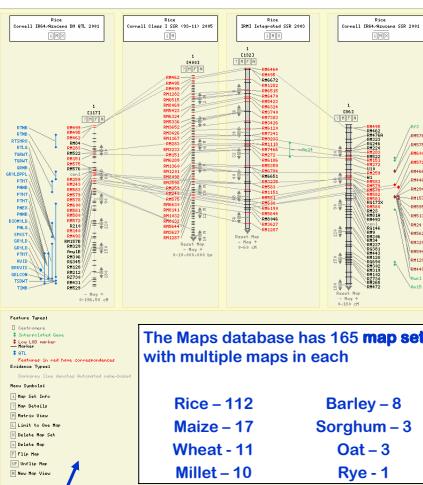


Customize the Genome Browser Detail View to show features of interest to you!

Browse the genome to find features, base pairs and codon options.

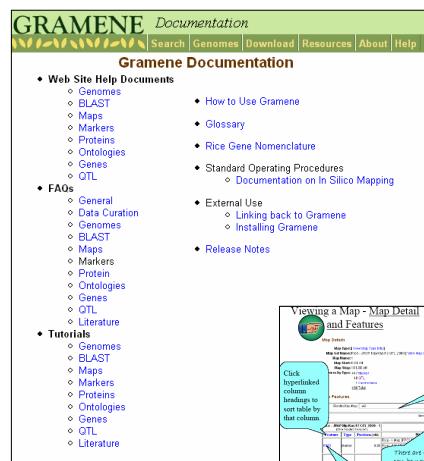


The Gramene website provides access to the Gramene databases and other online resources.



Rice – 112      Barley – 8  
Maize – 17      Sorghum – 3  
Wheat – 11      Oat – 3  
Millet – 10      Rye – 1

Use the Genome Browser to view features from several different maps in a single detailed view.



Genetic maps offer an indirect estimate of the distance and order of the sequence, and they provide the framework for genome sequencing.

Physical maps provide an estimate of the true distances (in base pairs), and define the sequence between markers. Used for gene identification.

Map Type: Genetic  
Accession ID: genetic  
Map Units: cM

Map Type: QTL  
Accession ID: qtl  
Map Units: cm

Map Type: Physical  
Map Units: bands  
Is Relational Only

Map Type: Deletion  
Accession ID: deletion  
Map Units: arm fraction

Map Type: Sequence  
Accession ID: sequence  
Map Units: bp

Online tutorials, FAQ's and other help documentation assist users in navigating and using the website. List-serve and feedback links allow users to communicate with database developers.

- PowerPoint (for non-IE)
- Adobe PDF
- Webpage (for IE, Safari, Konqueror)