

Resource Summary Report

Generated by [dkNET](#) on Aug 17, 2026

Plant Genomes Central

RRID:SCR_006541

Type: Tool

Proper Citation

Plant Genomes Central (RRID:SCR_006541)

Resource Information

URL: <http://www.ncbi.nlm.nih.gov/genomes/PLANTS/PGC-word.pdf>

Proper Citation: Plant Genomes Central (RRID:SCR_006541)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented August 19, 2016. A database of completed or in-progress sequenced plant genomes. The list of plant sequencing projects in this page includes those that have reached the stage where active sequence determination is currently producing, or is expected to produce in the near future. In addition, GenBank accession are provided toward the goal of determining the sequence of that plant genome.

Synonyms: Plant Genomes Central

Resource Type: data or information resource, database

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Plant Genomes Central

Resource ID: SCR_006541

Alternate IDs: nif-0000-03298

Old URLs: <http://www.ncbi.nlm.nih.gov/genomes/PLANTS/PlantList.html>

Record Creation Time: 20220129T080236+0000

Record Last Update: 20260815T182851+0000

Ratings and Alerts

No rating or validation information has been found for Plant Genomes Central.

No alerts have been found for Plant Genomes Central.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

Listed below are recent publications. The full list is available at [dkNET](#) .

Chen X, et al. (2021) Tandem UGT71B5s Catalyze Lignan Glycosylation in *Isatis indigotica* With Substrates Promiscuity. *Frontiers in plant science*, 12, 637695.

Marchal C, et al. (2020) Comparative Genomics and Functional Studies of Wheat BED-NLR Loci. *Genes*, 11(12).

Kalendar R, et al. (2020) Long Tandem Arrays of Cassandra Retroelements and Their Role in Genome Dynamics in Plants. *International journal of molecular sciences*, 21(8).

Sprink T, et al. (2014) The splicing fate of plant SPO11 genes. *Frontiers in plant science*, 5, 214.

Chien TY, et al. (2013) Analysis of new functional profiles of protein isoforms yielded by ds exonization in rice. *Evolutionary bioinformatics online*, 9, 417.

Shangguan L, et al. (2013) Evaluation of genome sequencing quality in selected plant species using expressed sequence tags. *PloS one*, 8(7), e69890.

Schatz MC, et al. (2012) Current challenges in de novo plant genome sequencing and assembly. *Genome biology*, 13(4), 243.

Liu LY, et al. (2012) Genome-wide survey of ds exonization to enrich transcriptomes and proteomes in plants. *Evolutionary bioinformatics online*, 8, 575.

Ellstrand NC, et al. (2010) Crops gone wild: evolution of weeds and invasives from domesticated ancestors. *Evolutionary applications*, 3(5-6), 494.

Tengs T, et al. (2009) Characterization of unknown genetic modifications using high throughput sequencing and computational subtraction. *BMC biotechnology*, 9, 87.