| Search NCBI | | | | | | |
|---------------|----------|--------|------|--------------------|---------------|--|
| NCBI Datasets | Taxonomy | Genome | Gene | Command-line tools | Documentation | |

Genome assembly Genome_assembly_40024.Col.1





⚠ Status: GenBank GCA_910591555.1 is suppressed

This record was removed at the submitter's request. Please contact update@ebi.ac.uk for further details.

| Submitted GenBank assembly GCA_910591555.1 (suppressed) | • |
|---------------------------------------------------------|---|
| Taxon Vitis vinifera (wine grape) | |
| WGS project CAJUXU01 | |
| Assembly type haploid | |
| Submitter INRAE Grand Est-Colmar | |
| Date Jul 7, 2021 | |



Assembly statistics

| | GenBank |
|-----------------------|----------|
| Genome size | 462.2 Mb |
| Total ungapped length | 460.5 Mb |
| Number of scaffolds | 19 |
| Scaffold N50 | 24.4 Mb |
| Scaffold L50 | 9 |

| | GenBank |
|-------------------|----------|
| Number of contigs | 2,177 |
| Contig N50 | 439.3 kb |
| Contig L50 | 319 |
| GC percent | 34.5 |
| Genome coverage | 30x |
| Assembly level | Scaffold |

Sample details

BioSample ID

SAMEA8898608

Description

PN40024

Comment

The accession of PN40024 with plant code 40024.Col.1 origins from nine rounds of self-pollination of Helfensteiner variety performed at INRAE Colmar, Haut-Rhin (68), France (WIEWS code: FRA038). This plant is grafted and available at INRAE Colmar, Haut-Rhin (68), France (WIEWS code: FRA038) in the genetic resource conservatory. This plant was used for extraction of genomic DNA for Pacific Biosciences sequencing for the update of the reference genome assembly PN40024.v4. Linked to this sample you can find the sequel run with raw pacbio data and two genome assembly (reference version (REF - haplotype 1) and alternative version (ALT - haplotype 2))

Submitter

EBI

ENA first public

Jun 7, 2021

ENA last update

Jan 13, 2022

ENA-CHECKLIST

ERC000011

View more ~

Assembly methods

Commen

This assembly, PN40024.v4, is the update of the previous Vitis vinifera reference genome assembly 12X.v2. It is not a de novo assembly, the reads was used to improve the 12X.v2 assembly (add of missing sequences, gap filling and scaffolding with PacBio reads). CHROMOSOME assembly.

Additional genomes

Browse all Vitis vinifera genomes (75)

BioProject

PRJEB45423

PN40024.v4 assembly

Chromosomes



Note: This scaffold-level genome assembly includes 19 scaffolds and no assembled chromosomes.

Revision history

This record has not been revised

| GenBank | RefSeq | Name | Level | Date | Action |
|-----------------|--------|----------------------|----------|-------------|--------|
| GCA_910591555.1 | n/a | Genome_assembly_4002 | Scaffold | Jul 7, 2021 | • |

FOLLOW NCBI

 \mathbb{X}

f

in



Connect with NLM



National Library of Medicine 8600 Rockville Pike Bethesda, MD 20894

Web Policies FOIA HHS Vulnerability Disclosure

Help Accessibility Careers

NLM NIH HHS USA.gov