

## GENOMES%0A

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[Genomes | Ensembl Genomes](#)

Match part of a name: elegans Wildeard .\* may be used: esch.\*coli; Division

[Scottish Genomes Partnership | Edinburgh | Glasgow ...](#)

The Scottish Genomes Partnership is a major Scotland-wide research programme between the Universities of Edinburgh, Glasgow, Aberdeen and Dundee, with NHS Scotland, NHS Lothian, NHS Greater Glasgow & Clyde, NHS Grampian and NHS Tayside.

[Ensembl Genomes](#)

Ensembl Genomes is developed by EMBL-EBI and is powered by Ensembl software system for the analysis and visualisation of genomic data. For details of our funding please click here.

[What's a Genome? - Genome News Network - Home GENOMES](#). A genome is all of a living thing's genetic material. It is the entire set of hereditary instructions for building, running, and maintaining an organism, and passing life on to the next generation.

[Genomes & Maps - Site Guide - NCBI](#)

Databases Assembly. A database providing information on the structure of assembled genomes, assembly names and other meta-data, statistical reports, and links to genomic sequence data.

[Full Genomes Corp](#)

Whole Genome Sequencing and Analysis. Our Whole Genome Sequencing and Analysis offers the most comprehensive genetic analysis available today. With 1000x more data than other commercial tests, results can be analyzed in exquisite detail.

[Silencing Genomes](#)

projects. DNA Learning Center Home MORE DNALC SITES: DNA from the Beginning DNA Interactive Eugenics Archive Genes to Cognition Online Your Genes, Your Health

[About | 1000 Genomes](#)

About IGSR and the 1000 Genomes Project. The 1000 Genomes Project ran between 2008 and 2015, creating the largest public catalogue of human variation and genotype data.

[1000 Genomes | A Deep Catalog of Human Genetic Variation](#)

IGSR and the 1000 Genomes Project. The International Genome Sample Resource (IGSR) was established to ensure the ongoing usability of data generated by the 1000 Genomes Project and to extend the data set.

[1001 Genomes](#)

The 1001 Genomes Vision. The 1001 Genomes Project

was launched at the beginning of 2008 to discover detailed whole-genome sequence variation in at least 1001 strains (accessions) of the reference plant *Arabidopsis thaliana*.

### **JGI IMG Integrated Microbial Genomes & Microbiomes**

The mission of the Integrated Microbial Genomes & Microbiomes (IMG/M) system is to support the annotation, analysis and distribution of microbial genome and microbiome datasets sequenced at DOE's Joint Genome Institute (JGI).

### **Submissions | Genome | Submission Portal**

Prokaryotic and eukaryotic genomes. Genomes is for complete, draft or incomplete genomes of prokaryotes or eukaryotes. Sequences should be at least 200 bp

### **100,000 whole genomes sequenced in the NHS - GOV.UK**

The 100,000 Genomes Project has sequenced 100,000 whole genomes, helping to improve diagnosis and treatment for patients with rare diseases and cancer.

### **Optimizer: A web server utility that optimize - Anglerfish**

OPTIMIZER is an on-line PHP application that optimizes the codon usage of a DNA sequence to increase its expression level. Users can introduce their own preference tables to be used in the optimization process or use pre-computed tables from more than 150 prokaryotic species under a strong translational selection.

### **In silico molecular biology experiments: PCR, multiplex ...**

Theoretical PCR amplification, AFLP-PCR and restriction digest of complete bacterial genomes, genomic data from sequenced prokaryotes.

### **Salk *Arabidopsis thaliana* 1,001 Genomes : Home Page**

*Arabidopsis* 2010 - 1,001 Genomes : Plant biologists need many completely sequenced and functionally annotated genomes within each species in order to fully exploit the power of evolution to understand how an organism functions and adapts to its environment.

### **Home - www.10wheatgenomes.com**

2017 was a remarkable year and represented a significant step forward for wheat genomics. In 2017, highly continuous genome assemblies were released for two of modern wheat's ancestors, wild emmer wheat (accession Zavitan ) 1 and *Aegilops tauschii* 2.

### **PaVE: Papilloma virus genome database**

Searchable database of papilloma genomes with visualization and analysis tools.

### **GtRNAdb: Genomic tRNA Database**

The genomic tRNA database contains tRNA gene predictions made by tRNAscan-SE on complete or nearly complete genomes. Unless otherwise noted, all annotation is automated, and has not been inspected for agreement with published literature.

#### **Xenbase Home**

Hello! We notice that you are using Internet Explorer, which is not supported by Xenbase and may cause the site to display incorrectly. We suggest using a current version of Chrome, FireFox, or Safari.