

## Search Ensembl

Search:  for



e.g. **human gene BRCA2** or **rat X:100000..200000** or **insulin**

## Browse a Genome

The Ensembl project produces genome databases for vertebrates and other eukaryotic species, and makes this information freely available online.

Click on a link below to go to the species' home page.

**Popular genomes** ([Log in to customize this list](#))



## All genomes

[View full list of all Ensembl species](#)

Other pre-build species are available in [Ensembl Pre!](#) →



Ensembl is a joint project between [EMBL - EBI](#) and the [Wellcome Trust Sanger Institute](#) to



develop a software system which produces and maintains automatic annotation on selected eukaryotic genomes.

Ensembl receives major funding from the Wellcome Trust. Our [acknowledgements page](#) includes a list of additional current and previous funding bodies.

## New to Ensembl?

Did you know you can:

[Add custom tracks](#)

using our new Control Panel

[Upload your own data](#)

and save it to your Ensembl account

[Search for a DNA or protein sequence](#)

using BLAST or BLAT

[Fetch only the data you want](#)

from our public database, using the Ensembl Perl API

[Download our databases via FTP](#)

in FASTA, MySQL and other formats

[Mine Ensembl with BioMart](#)

and export sequences or tables in text, html, or Excel format

Still got questions? [Try our FAQs](#)



### The new Ensembl website

We've made some changes to our site, to make it faster and easier to use.

[Find out more about what we've changed and why!](#)

## What's New in Release 51 (18 November 2008)

- [Webcode version 4.0 released](#) (all species)
- [New 2x genomes](#) (multiple species)
- [New Guineapig assembly and genebuild](#) (Guinea Pig)
- [Human patch for 51](#) (Human)
- [New xref sources](#) (Human)

[More news...](#)

## Latest Blog Entries

- Wed, 19 Nov 2008: [Upcoming training events December](#)
- Mon, 17 Nov 2008: [Accessing the Ensembl data with Perl](#)
- Fri, 31 Oct 2008: [Ensembl at the IMGIC](#)

[Go to Ensembl blog](#) →