

Resource Summary Report

Generated by [ASWG](#) on Aug 4, 2026

Genomicus

RRID:SCR_011791

Type: Tool

Proper Citation

Genomicus (RRID:SCR_011791)

Resource Information

URL: <http://www.genomicus.biologie.ens.fr/genomicus-72.01/cgi-bin/search.pl>

Proper Citation: Genomicus (RRID:SCR_011791)

Description: A genome browser that enables users to navigate in genomes in several dimensions: linearly along chromosome axes, transversally across different species, and chronologically along evolutionary time.

Abbreviations: Genomicus

Resource Type: data or information resource, database

Defining Citation: [PMID:23193262](#)

Keywords: genome, gene, synteny, browser, FASEB list

Resource Name: Genomicus

Resource ID: SCR_011791

Alternate IDs: OMICS_00914

Record Creation Time: 20220129T080306+0000

Record Last Update: 20260804T043453+0000

Ratings and Alerts

No rating or validation information has been found for Genomicus.

No alerts have been found for Genomicus.

Data and Source Information

Usage and Citation Metrics

We found 52 mentions in open access literature.

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

Qin X, et al. (2025) Goldfish phoenixin: (I) structural characterization, tissue distribution, and novel function as a feedforward signal for feeding-induced food intake in fish model. *Frontiers in endocrinology*, 16, 1570716.

Vélez EJ, et al. (2025) Endosomal microautophagy is activated by specific cellular stresses in trout hepatocytes. *Scientific reports*, 15(1), 39347.

Allio R, et al. (2025) Transcriptomic Data Reveal Divergent Paths of Chitinase Evolution Underlying Dietary Convergence in Anteaters and Pangolins. *Genome biology and evolution*, 17(2).

Merello Oyarzún G, et al. (2025) Evolutionary and Functional Analysis of Monoamine Oxidase F: A Novel Member of the Monoamine Oxidase Gene Family. *Genome biology and evolution*, 17(2).

Chen B, et al. (2025) Functional analysis of key members affecting egg production in the transglutaminase gene family in chickens. *Poultry science*, 104(2), 104794.

Ferez-Puche M, et al. (2024) Identification of a Novel γ -Defensin Gene in Gilthead Seabream (*Sparus aurata*). *Marine biotechnology* (New York, N.Y.), 26(6), 1219.

Roy J, et al. (2023) Characterization of free fatty acid receptor family in rainbow trout (*Oncorhynchus mykiss*): towards a better understanding of their involvement in fatty acid signalisation. *BMC genomics*, 24(1), 130.

Losilla M, et al. (2023) Molecular evolution of the ependymin-related gene *epdl2* in African weakly electric fish. *G3* (Bethesda, Md.), 13(3).

Zheng Z, et al. (2023) Phylogenetic and expression analysis of the angiopoietin-like gene family and their role in lipid metabolism in pigs. *Animal bioscience*, 36(10), 1517.

Gamboa M, et al. (2023) Evolutionary mechanisms underlying the diversification of nuclear factor of activated T cells across vertebrates. *Scientific reports*, 13(1), 6468.

Cardona E, et al. (2022) Physical Enrichment Triggers Brain Plasticity and Influences Blood Plasma Circulating miRNA in Rainbow Trout (*Oncorhynchus mykiss*). *Biology*, 11(8).

Opazo JC, et al. (2022) Identification of multiple TAR DNA binding protein retropseudogene lineages during the evolution of primates. *Scientific reports*, 12(1), 3823.

Parey E, et al. (2022) An atlas of fish genome evolution reveals delayed rediploidization following the teleost whole-genome duplication. *Genome research*, 32(9), 1685.

Xu Y, et al. (2022) Chicken CDS2 isoforms presented distinct spatio-temporal expression pattern and regulated by insulin in a breed-specific manner. *Poultry science*, 101(6), 101893.

Chen Y, et al. (2022) C/EBPZ modulates the differentiation and proliferation of preadipocytes. *International journal of obesity* (2005), 46(3), 523.

Xue T, et al. (2021) Genome-wide identification of interleukin-17 (IL-17) / interleukin-17 receptor (IL- 17R) in turbot (*Scophthalmus maximus*) and expression pattern analysis after *Vibrio anguillarum* infection. *Developmental and comparative immunology*, 121, 104070.

Noia M, et al. (2021) Characterization of the turbot *Scophthalmus maximus* (L.) myeloperoxidase. An insight into the evolution of vertebrate peroxidases. *Developmental and comparative immunology*, 118, 103993.

Ellingsen S, et al. (2021) The zebrafish cationic amino acid transporter/glycoprotein-associated family: sequence and spatiotemporal distribution during development of the transport system b0,+ (slc3a1/slc7a9). *Fish physiology and biochemistry*, 47(5), 1507.

Coppola U, et al. (2021) Origin and evolutionary landscape of Nr2f transcription factors across Metazoa. *PloS one*, 16(11), e0254282.

Boudinot P, et al. (2021) The repertoire of vertebrate STAT transcription factors: Origin and variations in fish. *Developmental and comparative immunology*, 116, 103929.