

Resource Summary Report

Generated by SPARC SAWG on Sep 18, 2026

Bioinformatics and Statistical Genetics

RRID:SCR_008589

Type: Tool

Proper Citation

Bioinformatics and Statistical Genetics (RRID:SCR_008589)

Resource Information

URL: <http://www.well.ox.ac.uk/bioinformatics-statistical-genetics>

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Description: Bioinformatics, statistical genetics, functional genomics, databases and scientific computing support and services for The Wellcome Trust Centre for Human Genetics to support the disease gene cloning projects within the centre and to maintain independent research in development of methods, algorithms and software for mapping multifactorial trait loci. The group has current research foci in the areas of sequence alignment statistics, comparative genomics, ancestral haplotype construction, LIMS database development, QTL linkage and association methods, variability in patterns of linkage disequilibrium and multivariate modelling of quantitative traits.

Abbreviations: WTCHG

Resource Type: data or information resource, portal, topical portal

Resource Name: Bioinformatics and Statistical Genetics

Resource ID: SCR_008589

Alternate IDs: nif-0000-31894

Record Creation Time: 20220129T080248+0000

Record Last Update: 20260912T195706+0000

Ratings and Alerts

No rating or validation information has been found for Bioinformatics and Statistical Genetics.

No alerts have been found for Bioinformatics and Statistical Genetics.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Poli M, et al. (2015) Characterization and quantification of proteins secreted by single human embryos prior to implantation. EMBO molecular medicine, 7(11), 1465.

Finlay C, et al. (2010) Chromosomal mapping of pancreatic islet morphological features and regulatory hormones in the spontaneously diabetic (Type 2) Goto-Kakizaki rat. Mammalian genome : official journal of the International Mammalian Genome Society, 21(9-10), 499.

Fearnside JF, et al. (2008) Phylometabonomic patterns of adaptation to high fat diet feeding in inbred mice. PloS one, 3(2), e1668.