

Resource Summary Report

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BETA

RRID:SCR_007556

Type: Tool

Proper Citation

BETA (RRID:SCR_007556)

Resource Information

URL: <http://cedar.genetics.soton.ac.uk/pub/PROGRAMS/BETA>

Proper Citation: BETA (RRID:SCR_007556)

Description: Software application for non-parametric linkage analysis using allele sharing in sib pairs (entry from Genetic Analysis Software)

Abbreviations: BETA

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c, unix, sun

Resource Name: BETA

Resource ID: SCR_007556

Alternate IDs: nlx_154241

Record Creation Time: 20220129T080242+0000

Record Last Update: 20260829T183049+0000

Ratings and Alerts

No rating or validation information has been found for BETA.

No alerts have been found for BETA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 138 mentions in open access literature.

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

Brooke GN, et al. (2026) Disruption of androgen receptor-cofactor interactions by the RNA-binding protein FUS/TLS alters androgen signalling in prostate cancer. *Oncogene*, 45(8), 757.

Shin S, et al. (2026) Generating multiple-choice questions using reverse engineering techniques. *Medical education online*, 31(1), 2611632.

Fahmy NM, et al. (2026) Artificial intelligence versus traditional approaches in multicomponent spectral analysis. *Scientific reports*, 16(1).

Sitohang IBS, et al. (2026) Comparative Profile of Microbiome in Normal Skin and Acne Vulgaris Skin Patients. *Clinical, cosmetic and investigational dermatology*, 19, 549168.

Schley RJ, et al. (2026) The frequency and importance of polyploidy in tropical rainforest tree radiations. *The New phytologist*, 249(4), 2103.

Shi H, et al. (2026) Effectiveness of nape acupuncture for post-stroke dysphagia: a meta-analysis and trial sequential analysis of randomized controlled trials. *Frontiers in neurology*, 17, 1720302.

Xiang Z, et al. (2026) 5-Hydroxymethylcytosine Dynamics Reveals Coordinated Reprogramming of Parental Genomes and X Chromosome Dosage Balance in Mouse SCNT Embryos. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 13(6), e09682.

Mei DA, et al. (2026) Liver disease in management and outcomes of European and Asian patients with atrial fibrillation: A report from two observational prospective registries. *European journal of clinical investigation*, 56(4), e70193.

Citu C, et al. (2025) Identification and catalog of viral transcriptional regulators in human diseases. *iScience*, 28(3), 112081.

Chambers TL, et al. (2025) At the Nexus Between Epigenetics and Senescence: The Effects of Senolytic (BI01) Administration on DNA Methylation Clock Age and the Methylome in Aged and Regenerated Skeletal Muscle. *Aging cell*, 24(7), e70068.

Cappello L, et al. (2025) Bayesian phylodynamic inference of population dynamics with dormancy. *Proceedings of the National Academy of Sciences of the United States of America*, 122(18), e2501394122.

Catozzi A, et al. (2025) Functional characterization of the ATOH1 molecular subtype indicates a pro-metastatic role in small cell lung cancer. *Cell reports*, 44(5), 115603.

Korn M, et al. (2025) SAVI Space-combinatorial encoding of the billion-size synthetically accessible virtual inventory. *Scientific data*, 12(1), 1064.

Urbanek A, et al. (2025) Molecular Determinants of Protein Pathogenicity at the Single-Aggregate Level. *Advanced science* (Weinheim, Baden-Wurttemberg, Germany), 12(9), e2410229.

Martinez-Marin D, et al. (2025) Dishevelled-1 regulates global transcriptomic changes and associates with ETS1 transcription factor. *Nature communications*, 16(1), 6288.

Shahid A, et al. (2025) S- and R-Carvedilol Prevent Benzo(a)pyrene-Induced Lung Carcinogenesis. *Thoracic cancer*, 16(12), e70109.

Elsakhawy OK, et al. (2025) Genome Mining Reveals a Sactipeptide Biosynthetic Cluster in *Staphylococcus pseudintermedius*. *Veterinary sciences*, 12(7).

Teng M, et al. (2025) Circular RMST cooperates with lineage-driving transcription factors to govern neuroendocrine transdifferentiation. *Cancer cell*, 43(5), 891.

Leung RWT, et al. (2025) CORN 2.0 - Condition Orientated Regulatory Networks 2.0. *Computational and structural biotechnology journal*, 27, 1518.

Chen L, et al. (2025) Molecular characterization of Coxsackievirus A2 isolated from hand, foot and mouth disease in eastern China. *Microbial genomics*, 11(9).