Resource Summary Report

Generated by kravitz2 on Sep 18, 2026

AutoMap

RRID:SCR_013095
Type: Tool

Proper Citation

AutoMap (RRID:SCR_013095)

Resource Information

URL: http://sourceforge.net/projects/ligmap/files/

Proper Citation: AutoMap (RRID:SCR_013095)

Description: A tool for structural biology and drug design.

Abbreviations: AutoMap

Resource Type: software resource

Resource Name: AutoMap

Resource ID: SCR_013095

Alternate IDs: OMICS_01596

Record Creation Time: 20220129T080314+0000

Record Last Update: 20260912T195759+0000

Ratings and Alerts

No rating or validation information has been found for AutoMap.

No alerts have been found for AutoMap.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 100 mentions in open access literature.

Listed below are recent publications. The full list is available at kravitz2 .

Han C, et al. (2026) Whole-exome sequencing for the genetic diagnosis of early-onset high myopia and associated hereditary eye disorders. BMC medical genomics, 19(1).

Owrang D, et al. (2026) Expansion of Molecular and Clinical Aspects of EPS8L2 (DFNB106)-Associated Hearing Loss Emphasizes a Potential Therapeutic Window. Molecular neurobiology, 63(1), 354.

Swetha J, et al. (2026) Genetic Diagnosis of Non-Syndromic Hearing Loss in South Indian Consanguineous Families Using Whole-Exome Sequencing. Medicina (Kaunas, Lithuania), 62(6).

de la Calle Arregui C, et al. (2026) Loss of heterozygosity exposes germline mutations in complex I and drives Warburg metabolism in oncocytic carcinoma of the thyroid. Science advances, 12(27), eaee5417.

Walker-Milne NL, et al. (2025) A novel use of Stereo Baited Remote Underwater Video and Drop-Down Video for biodiversity and marine landscape mapping and prediction. PloS one, 20(4), e0319355.

Han H, et al. (2025) Exome sequencing of 18,994 ethnically diverse patients with suspected rare Mendelian disorders. NPJ genomic medicine, 10(1), 6.

Cauti FM, et al. (2025) Ventricular Duration Map Area as a Valuable and Effective Target for VT Ablation End Point in Ischemic Cardiomyopathy: The VEDUM FREEDOM Study. Circulation. Arrhythmia and electrophysiology, 18(11), e013867.

Corral-Serrano JC, et al. (2025) A novel recurrent ARL3 variant c.209G > A p.(Gly70Glu) causes variable non-syndromic dominant retinal dystrophy with defective lipidated protein transport in human retinal stem cell models. Human molecular genetics, 34(9), 821.

Ahmad R, et al. (2025) Identification of a Novel GRM1 Frameshift Variant in Two Pakistani Families Broadens the Genetic Landscape of Ultra-Rare Spinocerebellar Ataxia Type 13. Cerebellum (London, England), 24(5), 145.

Huang YS, et al. (2025) High-density mapping in catheter ablation for atrial fibrillation in Asia Pacific region: An observational study. Journal of arrhythmia, 41(1), e13168.

Lee H, et al. (2025) A Korean Patient With Leber Congenital Amaurosis and a Homozygous RPE65 Variant Originating From a Paternal Uniparental Isodisomy. Molecular genetics & genomic medicine, 13(1), e70060.

Rashid A, et al. (2025) Exome sequencing identifies a homozygous splice site variant in RP1 as the underlying cause of autosomal recessive retinitis pigmentosa in a Pakistani family. Annals of medicine, 57(1), 2470953.

Neupane A, et al. (2025) Genetic improvement of FHB and DON resistance by combining the Fhb1 gene with additional resistance QTL in winter wheat population. The plant genome, 18(3), e70084.

Monfrini E, et al. (2025) RRP12 Variants Are Associated With Autosomal Recessive Brain Calcifications. Movement disorders : official journal of the Movement Disorder Society, 40(12), 2792.

Lee S, et al. (2025) Clinical utility of genome sequencing in rare diseases: lessons from a single-center study of 1,452 Korean families. NPJ genomic medicine, 11(1), 2.

Ecker C, et al. (2025) Transcriptomic decoding of surface-based imaging phenotypes and its application to pharmacotranscriptomics. Nature communications, 16(1), 6727.

Ahmad R, et al. (2025) Rare autosomal recessive hereditary sensory and autonomic neuropathy type VI in a Pakistani family caused by a novel DST variant. Neurological sciences : official journal of the Italian Neurological Society and of the Italian Society of Clinical Neurophysiology, 46(12), 6815.

Dominik N, et al. (2025) Biallelic variants in ARHGAP19 cause a progressive inherited motor-predominant neuropathy. The Journal of clinical investigation, 135(23).

Nadeali Z, et al. (2025) Investigating TSHR gene variants in consanguineous families: novel insights into variable expression in familial congenital hypothyroidism. Frontiers in endocrinology, 16, 1559281.

Ahmad R, et al. (2025) QARS1 associated developmental epileptic encephalopathy: first report of a rare homozygous missense variant from Pakistan causing nonepileptic phenotype in a family of seven patients and a comprehensive review of the literature. Molecular biology reports, 52(1), 528.