

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 10, 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: 20260808T190019+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 [PRECISE-TBI](#) .

Owring 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.

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).

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), eade5417.

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).

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.

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.

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.