

Resource Summary Report

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

Roadmap

RRID:SCR_017207

Type: Tool

Proper Citation

Roadmap (RRID:SCR_017207)

Resource Information

URL: http://www.sb.fsu.edu/~rsref/Distribution/roadmap_distribution.htm

Proper Citation: Roadmap (RRID:SCR_017207)

Description: Software tool to display surface of macromolecule and its properties. Uses projections to map van der Waals or solvent accessible surface of macromolecule onto plane., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Resource Type: software resource, software application

Defining Citation: [PMID:8384042](#)

Keywords: Display, surface, macromolecule, property, projection, van der Waal

Funding: NSF ;

NIH ;

Medical Research Council

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Roadmap

Resource ID: SCR_017207

License: Purdue Research Foundation

License URLs: <http://www.sb.fsu.edu/~rsref/Distribution/Roadmap/license.pdf>

Record Creation Time: 20220129T080334+0000

Record Last Update: 20260801T191100+0000

Ratings and Alerts

No rating or validation information has been found for Roadmap.

No alerts have been found for Roadmap.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 14 mentions in open access literature.

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

Wang TM, et al. (2025) A genome-wide association study integrated with single-cell and bulk profiles uncovers susceptibility genes for nasopharyngeal carcinoma involved in tumorigenesis via regulation of T cells. *Genome biology*, 26(1), 195.

Liu Y, et al. (2024) Wemics: A Single-Base Resolution Methylation Quantification Method for Enhanced Prediction of Epigenetic Regulation. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 11(21), e2308884.

Caparso C, et al. (2023) Mobile Technology to Monitor and Support Health and Well-Being: Qualitative Study of Perspectives and Design Suggestions From Patients Undergoing Hematopoietic Cell Transplantation. *JMIR formative research*, 7, e49806.

Lynall ME, et al. (2022) Genetic variants associated with psychiatric disorders are enriched at epigenetically active sites in lymphoid cells. *Nature communications*, 13(1), 6102.

Gao L, et al. (2022) Potential mechanisms and prognostic model of eRNAs-regulated genes in stomach adenocarcinoma. *Scientific reports*, 12(1), 16545.

Osman N, et al. (2022) Exploring the effects of genetic variation on gene regulation in cancer in the context of 3D genome structure. *BMC genomic data*, 23(1), 13.

Gilley KN, et al. (2022) Risk Factors for COVID-19 in College Students Identified by Physical, Mental, and Social Health Reported During the Fall 2020 Semester: Observational Study Using the Roadmap App and Fitbit Wearable Sensors. *JMIR mental health*, 9(2), e34645.

Cislo C, et al. (2021) Monitoring beliefs and physiological measures in students at risk for COVID-19 using wearable sensors and smartphone technology: Protocol for a mobile health study. *JMIR research protocols*, 10(6).

Liu RK, et al. (2021) Identification of novel functional CpG-SNPs associated with Type 2 diabetes and birth weight. *Aging*, 13(7), 10619.

Rusch A, et al. (2021) A Roadmap to Inform the Implementation of Evidence-Based Collaborative Care Interventions in Communities: Insights From the Michigan Mental

Health Integration Partnership. *Frontiers in public health*, 9, 655999.

Chen H, et al. (2020) New insights on human essential genes based on integrated analysis and the construction of the HEGIAP web-based platform. *Briefings in bioinformatics*, 21(4), 1397.

Kumaran M, et al. (2020) Fine-mapping of a novel premenopausal breast cancer susceptibility locus at Chr4q31.22 in Caucasian women and validation in African and Chinese women. *International journal of cancer*, 146(5), 1219.

Dormann C, et al. (2020) Evaluation of a game-based training course to build capacity for insecticide resistance management in vector control programmes. *PloS one*, 15(10), e0240514.

Meyer NL, et al. (2019) Structure of the gene therapy vector, adeno-associated virus with its cell receptor, AAVR. *eLife*, 8.