

Resource Summary Report

Generated by [kravitz2](#) on Jul 29, 2026

UpSetPlot

RRID:SCR_023225

Type: Tool

Proper Citation

UpSetPlot (RRID:SCR_023225)

Resource Information

URL: <https://upsetplot.readthedocs.io/en/stable/>

Proper Citation: UpSetPlot (RRID:SCR_023225)

Description: Software Python implementation of UpSet plots to visualize set overlaps.

Resource Type: data visualization software, software resource, software application, data processing software

Defining Citation: [PMID:26356912](#)

Keywords: UpSet plots, Python, visualize set overlaps,

Funding: Austrian Science Fund ;
Air Force Research Laboratory ;
DARPA ;
NHGRI K99 HG007583

Availability: Free, Available for download, Freely available

Resource Name: UpSetPlot

Resource ID: SCR_023225

Record Creation Time: 20230204T050200+0000

Record Last Update: 20260729T044552+0000

Ratings and Alerts

No rating or validation information has been found for UpSetPlot.

No alerts have been found for UpSetPlot.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Burankova Y, et al. (2025) Privacy-preserving multicenter differential protein abundance analysis with FedProt. Nature computational science, 5(8), 675.

Tingvall-Gustafsson J, et al. (2025) Chromatin interaction-based annotation of regulatory elements reveals dynamic promoter-enhancer interactions in lymphocyte development. iScience, 28(7), 112855.

Wei G, et al. (2024) Comprehensive data mining reveals RTK/RAS signaling pathway as a promoter of prostate cancer lineage plasticity through transcription factors and CNV. Scientific reports, 14(1), 11688.

Chiaro J, et al. (2023) Development of mesothelioma-specific oncolytic immunotherapy enabled by immunopeptidomics of murine and human mesothelioma tumors. Nature communications, 14(1), 7056.

S??raudie I, et al. (2023) A new scaffold-free tumoroid model provides a robust preclinical tool to investigate invasion and drug response in Renal Cell Carcinoma. Cell death & disease, 14(9), 622.

Loureiro C, et al. (2022) Comparative Metagenomic Analysis of Biosynthetic Diversity across Sponge Microbiomes Highlights Metabolic Novelty, Conservation, and Diversification. mSystems, 7(4), e0035722.