## **Resource Summary Report**

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# ggrepel

RRID:SCR\_017393 Type: Tool

**Proper Citation** 

ggrepel (RRID:SCR\_017393)

#### **Resource Information**

URL: https://cran.r-project.org/package=ggrepel

Proper Citation: ggrepel (RRID:SCR\_017393)

**Description:** Software tool to provide text and label geoms for ggplot2 that help to avoid overlapping text labels. Labels repel away from each other and away from data points.

Resource Type: software resource, standalone software, software application

Keywords: text, label, geoms, avoid, overlapping

Funding:

Availability: Free, Available for download, Freely available

Resource Name: ggrepel

Resource ID: SCR\_017393

Alternate URLs: https://github.com/slowkow/ggrepel

License: GNU GPL v3

Record Creation Time: 20220129T080335+0000

Record Last Update: 20250429T055912+0000

**Ratings and Alerts** 

No rating or validation information has been found for ggrepel.

No alerts have been found for ggrepel.

### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 38 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>NIF</u>.

Fuchs Wightman F, et al. (2024) Influence of RNA circularity on Target RNA-Directed MicroRNA Degradation. Nucleic acids research, 52(6), 3358.

Zaychikova M, et al. (2024) Vic9 mycobacteriophage: the first subcluster B2 phage isolated in Russia. Frontiers in microbiology, 15, 1513081.

Hoyer MJ, et al. (2024) Combinatorial selective ER-phagy remodels the ER during neurogenesis. Nature cell biology, 26(3), 378.

Chan M, et al. (2024) A kinase to cytokine explorer to identify molecular regulators and potential therapeutic opportunities. eLife, 12.

Reheman A, et al. (2024) The Role of Rv1476 in Regulating Stress Response and Intracellular Survival of Mycobacterium tuberculosis. Current issues in molecular biology, 46(2), 1556.

Wang J, et al. (2024) Protocol for identifying and comparing molecular prognosis subtypes of IgAN using R. STAR protocols, 5(3), 103138.

Ramos-Levi AM, et al. (2024) Maternal genomic profile, gestational diabetes control, and Mediterranean diet to prevent low birth weight. iScience, 27(12), 111376.

Bénéjam J, et al. (2024) Phenotyping data coupled with RNA sequencing of apple genotypes exhibiting contrasted quantitative trait loci architecture for apple scab (Venturia inaequalis) resistance. Data in brief, 56, 110778.

de Wit S, et al. (2024) Heart failure-induced microbial dysbiosis contributes to colonic tumour formation in mice. Cardiovascular research, 120(6), 612.

Lin CP, et al. (2024) Environmental DNA-based biodiversity profiling along the Houdong River in north-eastern Taiwan. Biodiversity data journal, 12, e116921.

Hodonsky CJ, et al. (2024) Multi-ancestry genetic analysis of gene regulation in coronary

arteries prioritizes disease risk loci. Cell genomics, 4(1), 100465.

Li H, et al. (2024) Protocol for mapping T cell activation using single-cell RNA-seq. STAR protocols, 5(4), 103409.

Blagojevic A, et al. (2024) Heat stress promotes Arabidopsis AGO1 phase separation and association with stress granule components. iScience, 27(3), 109151.

Baldaccini M, et al. (2024) The helicase domain of human Dicer prevents RNAi-independent activation of antiviral and inflammatory pathways. The EMBO journal, 43(5), 806.

Mangilet AF, et al. (2024) The Arabidopsis U1 snRNP regulates mRNA 3'-end processing. Nature plants, 10(10), 1514.

Lehle JD, et al. (2024) An in vitro approach reveals molecular mechanisms underlying endocrine disruptor-induced epimutagenesis. eLife, 13.

Borgmästars E, et al. (2024) Metabolomics for early pancreatic cancer detection in plasma samples from a Swedish prospective population-based biobank. Journal of gastrointestinal oncology, 15(2), 755.

Younginger BS, et al. (2023) Enrichment of oral-derived bacteria in inflamed colorectal tumors and distinct associations of Fusobacterium in the mesenchymal subtype. Cell reports. Medicine, 4(2), 100920.

Thompson LJ, et al. (2023) Contrasting effects of fungicide and herbicide active ingredients and their formulations on bumblebee learning and behaviour. The Journal of experimental biology, 226(6).

Spari D, et al. (2023) Intestinal dysbiosis as an intraoperative predictor of septic complications: evidence from human surgical cohorts and preclinical models of peritoneal sepsis. Scientific reports, 13(1), 22921.